PCT

WORLD INTELLECTUAL PROPERTY ORGANIZATION International Bureau



INTERNATIONAL APPLICATION PUBLISHED UNDER THE PATENT COOPERATION TREATY (PCT)

(51) International Pater C12Q 1/68, C12 15/86	nt Classification 7: 2N 15/00, 15/09, 15/63,	A1	 (11) International Publication Number: WO 00/55375 (43) International Publication Date: 21 September 2000 (21.09.00)
(21) International Appli	ication Number: PCT/US0	0/072	(81) Designated States: AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, CA, CH, CN, CU, CZ, DE, DK, EE, ES, FI, GB,
(22) International Filing Date: 17 March 2000 (17.03.00)			
			MW, MX, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK,
(30) Priority Data:			SL, TJ, TM, TR, TT, UA, UG, UZ, VN, YU, ZW, ARIPO
60/124,916	17 March 1999 (17.03.99)	τ	patent (GH, GM, KE, LS, MW, SD, SL, SZ, TZ, UG, ZW),
60/124,808	17 March 1999 (17.03.99)		Eurasian patent (AM, AZ, BY, KG, KZ, MD, RU, TJ, TM),
60/149,639	17 August 1999 (17.08.99)	τ	European patent (AT, BE, CH, CY, DE, DK, ES, FI, FR,
60/157,247	1 October 1999 (01.10.99)	Į	GB, GR, IE, IT, LU, MC, NL, PT, SE), OAPI patent (BF,
60/167,824	29 November 1999 (29.11.99)) (BJ, CF, CG, CI, CM, GA, GN, GW, ML, MR, NE, SN,
60/182,711	15 February 2000 (15.02.00)	Į	IS TD, TG).

- (71) Applicant: ALPHAGENE, INC. [US/US]; 260 mings Park, Woburn, MA 01801 (US).
- (72) Inventors: VALENZUELA, Dario; 1081 Hill Road, Boxborough, MA 01719-1010 (US). YUAN, Olive; 292 Mystic Street, Arlington, MA 02174 (US). HOFFMAN, Heidi; 90 Houghton Mill Road, Lunenburg, MA 01462 (US). HALL, Jeff; 4 Alderwood Drive, Stratham, NH 03885 (US). RAPIEJKO, Peter, 63 Old Grafton Road, Upton, MA 01568 (US).
- (74) Agent: SPRUNGER, Suzanne, A.; American Home Products Corporation, Patent & Trademark Dept. - 2B, One Campus Drive, Parsippany, NJ 07054 (US).

With international search report.

Before the expiration of the time limit for amending the claims and to be republished in the event of the receipt of amendments.

- (54) Title: SECRETED PROTEINS AND POLYNUCLEOTIDES ENCODING THEM
- (57) Abstract

Novel polynucleotides and the proteins encoded thereby are disclosed.

FOR THE PURPOSES OF INFORMATION ONLY

Codes used to identify States party to the PCT on the front pages of pamphlets publishing international applications under the PCT.

AL.	Albania	ES	Spain	LS	, Lesotho	SI	Slovenia
AM	Armenia	FI	Finland	LT	Lithuania	SK ·	Slovakia
AT	Austria	FR	France	LU	Luxembourg	SN	Senegal
AU	Australia	GA	Gabon	LV	Latvia	SZ	Swaziland
AZ	Azerbaijan	GB	United Kingdom	MC	Monaco	TD	Chad
BA	Bosnia and Herzegovina	GE	Georgia	MD	Republic of Moldova	TG	Togo
BB	Barbados	GH	Ghana	MG	Madagascar	TJ.	Tajikistan
BE	Belgium	GN	Guinea	MK	The former Yugoslav	TM	Turkmenistan
BF	Burkina Faso	GR	Greece		Republic of Macedonia	TR	Turkey
BG	Bulgaria	HU	Hungary	ML	Mali	TT	Trinidad and Tobago
ВЈ	Benin	IE	Ireland	MN	Mongolia	UA	Ukraine
BR	Brazil	IL	Israel	MR	Mauritania	UG	Uganda
BY	Belarus	IS	Iceland	MW	Malawi	US	United States of America
CA	Canada	IT	Italy .	MX	Mexico	UZ	Uzbekistan
CF	Central African Republic	JP	Japan	NE	Niger	VN	Vict Nam
CG	Congo	KE	Kenya	NL	Netherlands	YU	Yugoslavia
CH	Switzerland	KG	Kyrgyzstan	NO	Norway	2W	Zimbabwe
CI	Côte d'Ivoire	KP	Democratic People's	. NZ	New Zealand		
CM	Cameroon		Republic of Korea	PL	Poland		
CN	China	KR	Republic of Korea	PT	Portugal		
CU	Cuba	KZ	Kazakstan	RO	Romania		•
CZ	Czech Republic	LC	Saint Lucia	RU	Russian Federation		
DE	Germany	LI	Liechtenstein	SD	Sudan		
DK	Denmark	LK	Sri Lanka	SE	Sweden		
EE	Estonia	LR	Liberia	SG	Singapore		

Asn Lys His Ser Arg Ala Cys Gln Gln Phe Leu Lys Gln Cys Gln Leu

105

100

Arg Ser Phe Ala Leu Pro Leu 115 <210> 85 <211> 1176 <212> DNA <213> Homo sapiens <400> 85 · aatcttttt taaataagta ttttgttgag gttgaagaat tgctggcaat taaaagaata 60 gagetaatta tggettteat catteattea tgtatttatt gageacetae ttattatggt 120 gctcaacact tgttactgca agctacctta atttcccaag agtggtgcct tactctgttt 180 . tttctgatat ggtcttccaa tcagtgtgtg taacatacct gttgtttatc agccattgta 240 ggtggctgtg tctgttgcat catcataaga agtttaagct ttgtgctctg ataaattgtg 300 ttetgttaaa gaggttagta ggatgaaaac agcaaaacaa taatttttc aacaaattgt 360 aaattataag aaaaagagtt ggtttgtgta caacaatttt aatgattccc ttgttcattr 420 ttgctgtgaa atgcactgaa aaaaatcctc aaaatgagtt atagttcctg tgttgggaaa 480 attgacaaat aataaaacta gagaacaaac aataatgctt ctgtctcttt tacgaatgga 540 gagagaaagt ttatattcag tagagttatt gccctgttca tttgagaggg gcatggattt 600 tetgtttaag teetteagga atetteaget aggtggtaaa tttaataaga gtttetaaaa 660 attgaaatgt ttaactttta aatattetgg agatagaaga agaatataaa atgaaaccag 720 gctgatctgc atgcagtggc atttacaact aactgatcac aaccaattat agattcctta 780 ttttgtttat tgtgaggcag agtctgactc tgtcacccag gatggagtac agtgcatagc 840 tractgragt cttgacctcc caggetaaac caatcattcc acttcaccct cccaagtage 900 tgagaccaca ggcacacaac accacaacca gctgattgtt gtactgtong on a good accacaacca gctgattgtt gtactgtong on a good accacaacca grandaga etggtettga attectgage teaageagte eteceacete 1020 agecteccaa agtactggga ttacaggegt gaggtacete gece______ <210> 86 <211> 78 <212> PRT <213> Homo sapiens <400> 86 Met Tyr Leu Leu Ser Thr Tyr Leu Leu Trp Cys Ser Thr Leu Val Thr 1 5 10 15 Ala Ser Tyr Leu Asn Phe Pro Arg Val Val Pro Tyr Ser Val Phe Ser 20 25 30 Asp Met Val Phe Gin Ser Val Cys Val Thr Tyr Leu Leu Phe Ile Ser 35 40 45 His Cys Arg Trp Leu Cys Leu Leu His His His Lys Lys Phe Lys Leu 55 Cys Ala Leu Ile Asn Cys Val Leu Leu Lys Arg Leu Val Gly **65** · 70 <210> 87 <211> 1476 <212> DNA <213> Homo sapiens

1176

.

PCT/US00/07285 WO 00/55375

```
<400> 87
gaggetgeag aggtgecatg getgagteae acctgetgea gtggetgetg etgetgetge 60
ccaegetetg tggcccagge actgctgcct ggaccacctc atecttggcc tgtgcccagg 120
gecetgagtt etggtgeeaa ageetggage aageattgea gtgeagagee etagggeatt 180
gectacagga agtetgggga catgtgggag eegatgaeet atgecaagag tgtgaggaea 240
tegtecacat cettaacaag atggecaagg aggecatttt ceaggatete teegageage 300
aatteceeat teeteteece tattgetgge etetgeaggg etetgateaa geggateeaa 360
gccatgattc ccaagggtgc gctagctgtg gcagtggccc aggtgtgccg cgtggtacct 420
etggtggegg geggeatetg ceagtgeetg getgageget acteegteat cetgetegae 480
acguigetigg geografiget geoccagety gittgeografic tegiteties gitgetecaty 540
gatgacageg etggeceaag agaatggetg eegegagaet etgagtgeea cetetgeatg 600
tecgtgacea cecaggeegg gaacageage gageaggeea taccacagge aatgetecag 660
gcctgtgttg gctcctggct ggacagggaa aagtgcaagc aatttgtgga gcagcacacg 720
ccccagctge tgaccctggt gcccagggge tgggatgccc acaccacctg ccaggccctc 780
ggggtgtgtg ggaccatgte cagecetete cagtgtatee acageceega cetttgatga 840
gaactcagct gtccagaaaa agacaccgtc ctttaaagtg ctgcagtatg gccagacgtg 900
gtggetcaca cetgcaatce cagcacetta ggaggeegag gcaggaggat cettgaggte 960
aggagttega gaecageete gecaacatgg tgaaaceeca tttetaetaa aaatacaaaa 1020
aattagccaa gtgtggtggc atatgcctgt aatcccaact actcagaagg ccgaggcagg 1080
agaattactt gaacgcagga gaatcactgc agcccaggag gcagaggttg cagtgagccg 1140
agattgcacc actgcactcc agcctgggtg acagagcaag actccatctc agtaaataaa 1200
taaataaata aaaagegetg cagtagetgt ggeetcacce tgaagteage gggeecagge 1260
ctaceteact etetecettg geagagaage agaegteeat ageteetete eeteacaage 1320
geteceagee tgeeeteeag etgetgetet ecceteceag tetetaetea etgggatgag 1380
gttaggtcat gaggacacca aaaacctaaa aataaacaaa aagccaaaca agccttagct 1440
tttcttaaag acaaaaaaaa aaaaaaaa aaaaaa
<210> 88
<211> 145
<212> PRT
<213> Homo sapiens
<400> 88
Met Ala Glu Ser His Leu Leu Gln Trp Leu Leu Leu Leu Pro Thr
  1
                                     10
                                                          15
Leu Cys Gly Pro Gly Thr Ala Ala Trp Thr Thr Ser Ser Leu Ala Cys
             20
                                 25
                                                      30
Ala Gln Gly Pro Glu Phe Trp Cys Gln Ser Leu Glu Gln Ala Leu Gln
         35
Cys Arg Ala Leu Gly His Cys Leu Gln Glu Val Trp Gly His Val Gly
Ala Asp Asp Leu Cys Gln Glu Cys Glu Asp Ile Val His Ile Leu Asn
Lys Met Ala Lys Glu Ala Ile Phe Gln Asp Leu Ser Glu Gln Gln Phe
                 85
Pro Ile Pro Leu Pro Tyr Cys Trp Pro Leu Gln Gly Ser Asp Gln Ala
                                105
                                                    110
Asp Pro Ser His Asp Ser Cln Cly Cys Ala Ser Cys Gly Ser Gly Pro
        115
                            120
Gly Val Pro Arg Cly Thr Ser Gly Gly Gly Arg His Leu Pro Val Pro
    130
                                           - 140
```

1476

```
Gly
145
<210> 89
<211> 2243
<212> DNA
<213> Homo sapiens
<400> 89 .
agtactttgt aatcaagtga aaatataact ttatttttta actctattac attttattt 60
gtcatgtact assattattt ctgtattgct tttatassas acagtggcat ttagcactgg 120
cattgagact atagcacatc atttttgcca ttttcagtgc ttatattgtt aggtagaggc 180
tggcacttta ttagaatgca agccacaaaa atatcaattt tgttttttt gttagggtgg 240
gtettetttt tttettteee tetetettt tttaacaaat geettettat agaaaaactt 300
tctaagaggc aacaatttag aatggatatt ttgacgaatc ggcatgagtg taacagtgat 360
aacctgatct gtttgtttta aagattatta ccaagtgaaa aattcagaat gaatagaatt 420
tacactaaca tgctatataa aatgttanag tctgatgctg tgaaagcaat ctagtgctat 480
atttctacct cctcatttgt cttaattatt tggtaagtgg gattatgatg agtaactgga 540
ggggcttaga aacaaaaact ggatgaaaga gtatgcatga agaaaagctt ctttgataaa 600
tgtggagttc ttcattataa atatatattc atgaattcac agataagtac ttaaagaaca 660
gacagtttac ttggcctaaa aatattttga tgtttactca aaaagtacct cttcaggtct 720
tgagaacatg gaaaagaatt gagtgctttt aaatactttt tagaaagtaa tcataaaagt 780
aaattgaatt tcaaacctat ttggcttctg ttttgtgaac ctttgaacta tatgtatgtg 840
tataagggta tacacataca tatatggcat ataacaagtg tacacatata cacataacaa 900
gtgtagaagt atatattaca tacatacact cactetgtet ggtatagget aattttgaag 960
aacteccata agtttetget getteteeca taactgetge caccaccate agaatteata 1020
atcaaaccta acctttttgt ttggggcacc aaatctgaag acaaaattaa tttgcaccag 1080
taaacttcaa getgetttet ttettgaaaa etaaaegttt aaegtataat gtetgtttgg 1140
atactgttcc aaattgttga ttgcatgtgg ttaatgttgc attagagcac tttgcaattg 1200
cataattcat taatgttttg tgagcttgca tttgtgagtt attggatgat cagactgaat 1260
tttgtcaagt atcacattgt acatcttgcc tagatgtcga tgactgcaag taataataca 1320
gtttataatg aaactatcta caattcttgt tttagcacat ctgttatccg taaaacacct 1380
gtaactaget tttttaattt attatttgaa ttttaggata gegaateact aatttttagt 1440
tgctgaggtt ggcattttag tgattattaa gcacttctgt cagtctttga aaaaagaacg 1500
tattttttgt gctttgaaga tctctgaaga atttctttta taatagaatg ggcatgtatt 1560
gtaacagttt tatgtcaaat gatctgtgct gtagaaaaac attaaccett gttcaaaaaa 1620
gaaatggata aacttggcct ttctaagtgg taagaatgac ctgtcactat aatatactgt 1680
atgtttacat tttatttaaa ttttatctct tatgtatagg gtgataacct tccccagaaa 1740
caacagtgat tgcgattgtt ttctagaaac ttctttaaag tgccacattt ggcagtacaa 1800
atgagtetga gtgtaatage eeagagattt atatatagtt gaatgtetaa aatggtaaaa 1860
tgtgccactg tgtcaagtta cagtggctta tgtttttcat agtaattcaa atgaacttcc 1920
tatttttgat agtaaatgtc atttaatagt atacttgcca tttgagcctc actgcaaaat 1980
tagtgcagag gagaaaacaa tttttaatgt aatcttgatt ttacctcata tactgtacat 2040
tccaaaaact ctaaactttt taaagattat agatacacta ccaaacatat caccttaaaa 2100
ttgtataagg ctgaatgaac ttcatacaaa tgaaaaaaat ctcataaaaa tacataaact 2160
atgtagcaaa agtatctgta aaatccatgg aaaataaaag ttgtatcatt ctttttgaaa 2220
                                                                   2243
aaa aaaaaaaa aaaaaaaaa aaa
<210> 90
<211> 61
<212> PRT
<213> Homo sapiens
<400> 90
Met Gln Ala Thr Lys Ile Ser Ile Leu Phe Phe Leu Leu Gly Trp Val
```

Phe Phe Phe Ser Phe Pro Leu Ser Phe Phe Asn Lys Cys Leu Leu Ile

What is claimed is:

1. An isolated polynucleotide comprising a nucleotide sequence selected from the group consisting of:

- (a) the nucleotide sequence of SEQ ID NO:1;
- (b) the nucleotide sequence of SEQ ID NO:1 from nucleotide 27 to nucleotide 260;
- (c) the nucleotide sequence of SEQ ID NO:1 from nucleotide 72 to nucleotide 260;
- (d) the nucleotide sequence of the full-length protein coding sequence of clone vc62_1 deposited with the ATCC under accession number 207114;
- (e) a nucleotide sequence encoding the full-length protein encoded by the cDNA insert of clone vc62_1 deposited with the ATCC under accession number 207114;
- (f) the nucleotide sequence of a mature protein coding sequence of clone vc62_1 deposited with the ATCC under accession number 207114;
- (g) a nucleotide sequence encoding a mature protein encoded by the cDNA insert of clone vc62_1 deposited with the ATCC under accession number 207114;
- (h) a nucleotide sequence encoding a protein comprising the amino acid sequence of SEQ ID NO:2;
- (i) a nucleotide sequence encoding a protein comprising a fragment of the amino acid sequence of SEQ ID NO:2, the fragment comprising eight contiguous amino acids of SEQ ID NO:2;
- (j) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 65 degrees C, or 4X SSC at 42 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g); and
- (k) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 50 degrees C, or 6X SSC at 40 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g), and that has a length that is at least 25% of the length of SEQ ID NO:1.

2. The polynucleotide of claim 1 wherein said polynucleotide is operably linked to at least one expression control sequence.

- 3. A host cell transformed with the polynucleotide of claim 2.
- 4. The host cell of claim 3, wherein said cell is a mammalian cell.
- 5. A process for producing a protein encoded by the polynucleotide of claim 2, which process comprises:
 - (a) growing a culture of a host cell in a suitable culture medium, wherein the host cell has been transformed with the polynucleotide of claim 2; and
 - (b) purifying said protein from the culture.
 - 6. A protein produced according to the process of claim 5.
 - 7. An isolated polynucleotide encoding the protein of claim 6.
- 8. The polynucleotide of claim 7, wherein the polynucleotide comprises the cDNA insert of clone vc62_1 deposited with the ATCC under accession number 207114.
- 9. A protein comprising an amino acid sequence selected from the group consisting of:
 - (a) the amino acid sequence of SEQ ID NO:2;
 - (b) a fragment of the amino acid sequence of SEQ ID NO:2, the fragment comprising eight contiguous amino acids of SEQ ID NO:2; and
- (c) the amino acid sequence encoded by the cDNA insert of clone vc62_1 deposited with the ATCC under accession number 207114; the protein being substantially free from other mammalian proteins.
- 10. The protein of claim 9, wherein said protein comprises the amino acid sequence of SEQ ID NO:2.

11. A composition comprising the protein of claim 9 and a pharmaceutically acceptable carrier.

- 12. An isolated polynucleotide comprising a nucleotide sequence selected from the group consisting of:
 - (a) the nucleotide sequence of SEQ ID NO:3;
 - (b) the nucleotide sequence of SEQ ID NO:3 from nucleotide 6 to nucleotide 1325;
 - (c) the nucleotide sequence of SEQ ID NO:3 from nucleotide 99 to nucleotide 1325;
 - (d) the nucleotide sequence of the full-length protein coding sequence of clone vp10_1 deposited with the ATCC under accession number 207114;
 - (e) a nucleotide sequence encoding the full-length protein encoded by the cDNA insert of clone vp10_1 deposited with the ATCC under accession number 207114;
 - (f) the nucleotide sequence of a mature protein coding sequence of clone vp10_1 deposited with the ATCC under accession number 207114;
 - (g) a nucleotide sequence encoding a mature protein encoded by the cDNA insert of clone vp10_1 deposited with the ATCC under accession number 207114;
 - (h) a nucleotide sequence encoding a protein comprising the amino acid sequence of SEQ ID NO:4;
 - (i) a nucleotide sequence encoding a protein comprising a fragment of the amino acid sequence of SEQ ID NO:4, the fragment comprising eight contiguous amino acids of SEQ ID NO:4;
 - (j) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 65 degrees C, or 4X SSC at 42 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g); and
 - (k) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 50 degrees C, or 6X SSC at 40 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g), and that has a length that is at least 25% of the length of SEQ ID NO:3.

13. A protein comprising an amino acid sequence selected from the group consisting of:

- (a) the amino acid sequence of SEQ ID NO:4;
- (b) a fragment of the amino acid sequence of SEQ ID NO:4, the fragment comprising eight contiguous amino acids of SEQ ID NO:4; and
- (c) the amino acid sequence encoded by the cDNA insert of clone vp10_1 deposited with the ATCC under accession number 207114; the protein being substantially free from other mammalian proteins.
- 14. An isolated polynucleotide comprising a nucleotide sequence selected from the group consisting of:
 - (a) the nucleotide sequence of SEQ ID NO:5;
 - (b) the nucleotide sequence of SEQ ID NO:5 from nucleotide 149 to nucleotide 322;
 - (c) the nucleotide sequence of SEQ ID NO:5 from nucleotide 200 to nucleotide 322;
 - (d) the nucleotide sequence of the full-length protein coding sequence of clone vp11_1 deposited with the ATCC under accession number 207114;
 - (e) a nucleotide sequence encoding the full-length protein encoded by the cDNA insert of clone vp11_1 deposited with the ATCC under accession number 207114;
 - (f) the nucleotide sequence of a mature protein coding sequence of clone vp11_1 deposited with the ATCC under accession number 207114;
 - (g) a nucleotide sequence encoding a mature protein encoded by the cDNA insert of clone vp11_1 deposited with the ATCC under accession number 207114;
 - (h) a nucleotide sequence encoding a protein comprising the amino acid sequence of SEQ ID NO:6;
 - (i) a nucleotide sequence encoding a protein comprising a fragment of the amino acid sequence of SEQ ID NO:6, the fragment comprising eight contiguous amino acids of SEQ ID NO:6;

(j) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 65 degrees C, or 4X SSC at 42 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g); and

- (k) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 50 degrees C, or 6X SSC at 40 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g), and that has a length that is at least 25% of the length of SEQ ID NO:5.
- 15. A protein comprising an amino acid sequence selected from the group consisting of:
 - (a) the amino acid sequence of SEQ ID NO:6;
 - (b) a fragment of the amino acid sequence of SEQ ID NO:6, the fragment comprising eight contiguous amino acids of SEQ ID NO:6; and
- (c) the amino acid sequence encoded by the cDNA insert of clone vp11_1 deposited with the ATCC under accession number 207114; the protein being substantially free from other mammalian proteins.
- 16. An isolated polynucleotide comprising a nucleotide sequence selected from the group consisting of:
 - (a) the nucleotide sequence of SEQ ID NO:7;
 - (b) the nucleotide sequence of SEQ ID NO:7 from nucleotide 288 to nucleotide 629;
 - (c) the nucleotide sequence of SEQ ID NO:7 from nucleotide 363 to nucleotide 629;
 - (d) the nucleotide sequence of the full-length protein coding sequence of clone vp13_1 deposited with the ATCC under accession number 207114;
 - (e) a nucleotide sequence encoding the full-length protein encoded by the cDNA insert of clone vp13_1 deposited with the ATCC under accession number 207114;
 - (f) the nucleotide sequence of a mature protein coding sequence of clone vp13_1 deposited with the ATCC under accession number 207114;

(g) a nucleotide sequence encoding a mature protein encoded by the cDNA insert of clone vp13_1 deposited with the ATCC under accession number 207114;

- (h) a nucleotide sequence encoding a protein comprising the amino acid sequence of SEQ ID NO:8;
- (i) a nucleotide sequence encoding a protein comprising a fragment of the amino acid sequence of SEQ ID NO:8, the fragment comprising eight contiguous amino acids of SEQ ID NO:8;
- (j) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 65 degrees C, or 4X SSC at 42 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g); and
- (k) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 50 degrees C, or 6X SSC at 40 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g), and that has a length that is at least 25% of the length of SEQ ID NO:7.
- 17. A protein comprising an amino acid sequence selected from the group consisting of:
 - (a) the amino acid sequence of SEQ ID NO:8;
 - (b) a fragment of the amino acid sequence of SEQ ID NO:8, the fragment comprising eight contiguous amino acids of SEQ ID NO:8; and
- (c) the amino acid sequence encoded by the cDNA insert of clone vp13_1 deposited with the ATCC under accession number 207114; the protein being substantially free from other mammalian proteins.
- 18. An isolated polynucleotide comprising a nucleotide sequence selected from the group consisting of:
 - (a) the nucleotide sequence of SEQ ID NO:9;
 - (b) the nucleotide sequence of SEQ ID NO:9 from nucleotide 11 to nucleotide 298;
 - (c) the nucleotide sequence of SEQ ID NO:9 from nucleotide 149 to nucleotide 298;

(d) the nucleotide sequence of the full-length protein coding sequence of clone vp16_1 deposited with the ATCC under accession number 207114;

- (e) a nucleotide sequence encoding the full-length protein encoded by the cDNA insert of clone vp16_1 deposited with the ATCC under accession number 207114;
- (f) the nucleotide sequence of a mature protein coding sequence of clone vp16_1 deposited with the ATCC under accession number 207114;
- (g) a nucleotide sequence encoding a mature protein encoded by the cDNA insert of clone vp16_1 deposited with the ATCC under accession number 207114;
- (h) a nucleotide sequence encoding a protein comprising the amino acid sequence of SEQ ID NO:10;
- (i) a nucleotide sequence encoding a protein comprising a fragment of the amino acid sequence of SEQ ID NO:10, the fragment comprising eight contiguous amino acids of SEQ ID NO:10;
- (j) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 65 degrees C, or 4X SSC at 42 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g); and
- (k) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 50 degrees C, or 6X SSC at 40 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g), and that has a length that is at least 25% of the length of SEQ ID NO:9.
- 19. A protein comprising an amino acid sequence selected from the group consisting of:
 - (a) the amino acid sequence of SEQ ID NO:10;
 - (b) a fragment of the amino acid sequence of SEQ ID NO:10, the fragment comprising eight contiguous amino acids of SEQ ID NO:10; and
- (c) the amino acid sequence encoded by the cDNA insert of clone vp16_1 deposited with the ATCC under accession number 207114; the protein being substantially free from other mammalian proteins.

20. An isolated polynucleotide comprising a nucleotide sequence selected from the group consisting of:

- (a) the nucleotide sequence of SEQ ID NO:11;
- (b) the nucleotide sequence of SEQ ID NO:11 from nucleotide 257 to nucleotide 607;
- (c) the nucleotide sequence of SEQ ID NO:11 from nucleotide 479 to nucleotide 607;
- (d) the nucleotide sequence of the full-length protein coding sequence of clone vp21_1 deposited with the ATCC under accession number 207114;
- (e) a nucleotide sequence encoding the full-length protein encoded by the cDNA insert of clone vp21_1 deposited with the ATCC under accession number 207114;
- (f) the nucleotide sequence of a mature protein coding sequence of clone vp21_1 deposited with the ATCC under accession number 207114;
- (g) a nucleotide sequence encoding a mature protein encoded by the cDNA insert of clone vp21_1 deposited with the ATCC under accession number 207114;
- (h) a nucleotide sequence encoding a protein comprising the amino acid sequence of SEQ ID NO:12;
- (i) a nucleotide sequence encoding a protein comprising a fragment of the amino acid sequence of SEQ ID NO:12, the fragment comprising eight contiguous amino acids of SEQ ID NO:12;
- (j) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 65 degrees C, or 4X SSC at 42 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g); and
- (k) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 50 degrees C, or 6X SSC at 40 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g), and that has a length that is at least 25% of the length of SEQ ID NO:11.
- 21. A protein comprising an amino acid sequence selected from the group consisting of:

- (a) the amino acid sequence of SEQ ID NO:12;
- (b) a fragment of the amino acid sequence of SEQ ID NO:12, the fragment comprising eight contiguous amino acids of SEQ ID NO:12; and
- (c) the amino acid sequence encoded by the cDNA insert of clone vp21_1 deposited with the ATCC under accession number 207114; the protein being substantially free from other mammalian proteins.
- 22. An isolated polynucleotide comprising a nucleotide sequence selected from the group consisting of:
 - (a) the nucleotide sequence of SEQ ID NO:13;
 - (b) the nucleotide sequence of SEQ ID NO:13 from nucleotide 163 to nucleotide 477;
 - (c) the nucleotide sequence of SEQ ID NO:13 from nucleotide 238 to nucleotide 477;
 - (d) the nucleotide sequence of the full-length protein coding sequence of clone vp22_1 deposited with the ATCC under accession number 207114;
 - (e) a nucleotide sequence encoding the full-length protein encoded by the cDNA insert of clone vp22_1 deposited with the ATCC under accession number 207114;
 - (f) the nucleotide sequence of a mature protein coding sequence of clone vp22_1 deposited with the ATCC under accession number 207114;
 - (g) a nucleotide sequence encoding a mature protein encoded by the cDNA insert of clone vp22_1 deposited with the ATCC under accession number 207114;
 - (h) a nucleotide sequence encoding a protein comprising the amino acid sequence of SEQ ID NO:14;
 - (i) a nucleotide sequence encoding a protein comprising a fragment of the amino acid sequence of SEQ ID NO:14, the fragment comprising eight contiguous amino acids of SEQ ID NO:14;
 - (j) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 65 degrees C, or 4X SSC at 42 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g); and

(k) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 50 degrees C, or 6X SSC at 40 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g), and that has a length that is at least 25% of the length of SEQ ID NO:13.

- 23. A protein comprising an amino acid sequence selected from the group consisting of:
 - (a) the amino acid sequence of SEQ ID NO:14;
 - (b) a fragment of the amino acid sequence of SEQ ID NO:14, the fragment comprising eight contiguous amino acids of SEQ ID NO:14; and
- (c) the amino acid sequence encoded by the cDNA insert of clone vp22_1 deposited with the ATCC under accession number 207114; the protein being substantially free from other mammalian proteins.
- 24. An isolated polynucleotide comprising a nucleotide sequence selected from the group consisting of:
 - (a) the nucleotide sequence of SEQ ID NO:15;
 - (b) the nucleotide sequence of SEQ ID NO:15 from nucleotide 58 to nucleotide 624;
 - (c) the nucleotide sequence of SEQ ID NO:15 from nucleotide 106 to nucleotide 624;
 - (d) the nucleotide sequence of the full-length protein coding sequence of clone vq2_1 deposited with the ATCC under accession number 207114;
 - (e) a nucleotide sequence encoding the full-length protein encoded by the cDNA insert of clone vq2_1 deposited with the ATCC under accession number 207114;
 - (f) the nucleotide sequence of a mature protein coding sequence of clone vq2_1 deposited with the ATCC under accession number 207114;
 - (g) a nucleotide sequence encoding a mature protein encoded by the cDNA insert of clone vq2_1 deposited with the ATCC under accession number 207114;

(h) a nucleotide sequence encoding a protein comprising the amino acid sequence of SEQ ID NO:16;

- (i) a nucleotide sequence encoding a protein comprising a fragment of the amino acid sequence of SEQ ID NO:16, the fragment comprising eight contiguous amino acids of SEQ ID NO:16;
- (j) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 65 degrees C, or 4X SSC at 42 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g); and
- (k) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 50 degrees C, or 6X SSC at 40 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g), and that has a length that is at least 25% of the length of SEQ ID NO:15.
- 25. A protein comprising an amino acid sequence selected from the group consisting of:
 - (a) the amino acid sequence of SEQ ID NO:16;
 - (b) a fragment of the amino acid sequence of SEQ ID NO:16, the fragment comprising eight contiguous amino acids of SEQ ID NO:16; and
- (c) the amino acid sequence encoded by the cDNA insert of clone vq2_1 deposited with the ATCC under accession number 207114; the protein being substantially free from other mammalian proteins.
- 26. An isolated polynucleotide comprising a nucleotide sequence selected from the group consisting of:
 - (a) the nucleotide sequence of SEQ ID NO:17;
 - (b) the nucleotide sequence of SEQ ID NO:17 from nucleotide 773 to nucleotide 1090;
 - (c) the nucleotide sequence of SEQ ID NO:17 from nucleotide 842 to nucleotide 1090;
 - (d) the nucleotide sequence of the full-length protein coding sequence of clone vq3_1 deposited with the ATCC under accession number 207114;

(e) a nucleotide sequence encoding the full-length protein encoded by the cDNA insert of clone vq3_1 deposited with the ATCC under accession number 207114;

- (f) the nucleotide sequence of a mature protein coding sequence of clone vq3_1 deposited with the ATCC under accession number 207114;
- (g) a nucleotide sequence encoding a mature protein encoded by the cDNA insert of clone vq3_1 deposited with the ATCC under accession number 207114;
- (h) a nucleotide sequence encoding a protein comprising the amino acid sequence of SEQ ID NO:18;
- (i) a nucleotide sequence encoding a protein comprising a fragment of the amino acid sequence of SEQ ID NO:18, the fragment comprising eight contiguous amino acids of SEQ ID NO:18;
- (j) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 65 degrees C, or 4X SSC at 42 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g); and
- (k) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 50 degrees C, or 6X SSC at 40 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g), and that has a length that is at least 25% of the length of SEQ ID NO:17.
- 27. A protein comprising an amino acid sequence selected from the group consisting of:
 - (a) the amino acid sequence of SEQ ID NO:18;
 - (b) a fragment of the amino acid sequence of SEQ ID NO:18, the fragment comprising eight contiguous amino acids of SEQ ID NO:18; and
- (c) the amino acid sequence encoded by the cDNA insert of clone vq3_1 deposited with the ATCC under accession number 207114; the protein being substantially free from other mammalian proteins.
- 28. An isolated polynucleotide comprising a nucleotide sequence selected from the group consisting of:

- (a) the nucleotide sequence of SEQ ID NO:19;
- (b) the nucleotide sequence of SEQ ID NO:19 from nucleotide 96 to nucleotide 275;
- (c) the nucleotide sequence of SEQ ID NO:19 from nucleotide 159 to nucleotide 275;
- (d) the nucleotide sequence of the full-length protein coding sequence of clone vq5_1 deposited with the ATCC under accession number 207114;
- (e) a nucleotide sequence encoding the full-length protein encoded by the cDNA insert of clone vq5_1 deposited with the ATCC under accession number 207114;
- (f) the nucleotide sequence of a mature protein coding sequence of clone vq5_1 deposited with the ATCC under accession number 207114;
- (g) a nucleotide sequence encoding a mature protein encoded by the cDNA insert of clone vq5_1 deposited with the ATCC under accession number 207114;
- (h) a nucleotide sequence encoding a protein comprising the amino acid sequence of SEQ ID NO:20;
- (i) a nucleotide sequence encoding a protein comprising a fragment of the amino acid sequence of SEQ ID NO:20, the fragment comprising eight contiguous amino acids of SEQ ID NO:20;
- (j) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 65 degrees C, or 4X SSC at 42 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g); and
- (k) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 50 degrees C, or 6X SSC at 40 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g), and that has a length that is at least 25% of the length of SEQ ID NO:19.
- 29. A protein comprising an amino acid sequence selected from the group consisting of:
 - (a) the amino acid sequence of SEQ ID NO:20;

(b) a fragment of the amino acid sequence of SEQ ID NO:20, the fragment comprising eight contiguous amino acids of SEQ ID NO:20; and

- (c) the amino acid sequence encoded by the cDNA insert of clone vq5_1 deposited with the ATCC under accession number 207114; the protein being substantially free from other mammalian proteins.
- 30. An isolated polynucleotide comprising a nucleotide sequence selected from the group consisting of:
 - (a) the nucleotide sequence of SEQ ID NO:21;
 - (b) the nucleotide sequence of SEQ ID NO:21 from nucleotide 176 to nucleotide 340;
 - (c) the nucleotide sequence of SEQ ID NO:21 from nucleotide 230 to nucleotide 340;
 - (d) the nucleotide sequence of the full-length protein coding sequence of clone vq6_1 deposited with the ATCC under accession number 207114;
 - (e) a nucleotide sequence encoding the full-length protein encoded by the cDNA insert of clone vq6_1 deposited with the ATCC under accession number 207114;
 - (f) the nucleotide sequence of a mature protein coding sequence of clone vq6_1 deposited with the ATCC under accession number 207114;
 - (g) a nucleotide sequence encoding a mature protein encoded by the cDNA insert of clone vq6_1 deposited with the ATCC under accession number 207114;
 - (h) a nucleotide sequence encoding a protein comprising the amino acid sequence of SEQ ID NO:22;
 - (i) a nucleotide sequence encoding a protein comprising a fragment of the amino acid sequence of SEQ ID NO:22, the fragment comprising eight contiguous amino acids of SEQ ID NO:22;
 - (j) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 65 degrees C, or 4X SSC at 42 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g); and

(k) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 50 degrees C, or 6X SSC at 40 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g), and that has a length that is at least 25% of the length of SEQ ID NO:21.

- 31. A protein comprising an amino acid sequence selected from the group consisting of:
 - (a) the amino acid sequence of SEQ ID NO:22;
 - (b) a fragment of the amino acid sequence of SEQ ID NO:22, the fragment comprising eight contiguous amino acids of SEQ ID NO:22; and
- (c) the amino acid sequence encoded by the cDNA insert of clone vq6_1 deposited with the ATCC under accession number 207114; the protein being substantially free from other mammalian proteins.
- 32. An isolated polynucleotide comprising a nucleotide sequence selected from the group consisting of:
 - (a) the nucleotide sequence of SEQ ID NO:23;
 - (b) the nucleotide sequence of SEQ ID NO:23 from nucleotide 29 to nucleotide 1111;
 - (c) the nucleotide sequence of SEQ ID NO:23 from nucleotide 167 to nucleotide 1111;
 - (d) the nucleotide sequence of the full-length protein coding sequence of clone vrl_1 deposited with the ATCC under accession number 207114;
 - (e) a nucleotide sequence encoding the full-length protein encoded by the cDNA insert of clone vrl_1 deposited with the ATCC under accession number 207114;
 - (f) the nucleotide sequence of a mature protein coding sequence of clone vr1_1 deposited with the ATCC under accession number 207114;
 - (g) a nucleotide sequence encoding a mature protein encoded by the cDNA insert of clone vrl_1 deposited with the ATCC under accession number 207114;

(h) a nucleotide sequence encoding a protein comprising the amino acid sequence of SEQ ID NO:24;

- (i) a nucleotide sequence encoding a protein comprising a fragment of the amino acid sequence of SEQ ID NO:24, the fragment comprising eight contiguous amino acids of SEQ ID NO:24;
- (j) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 65 degrees C, or 4X SSC at 42 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g); and
- (k) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 50 degrees C, or 6X SSC at 40 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g), and that has a length that is at least 25% of the length of SEQ ID NO:23.
- 33. A protein comprising an amino acid sequence selected from the group consisting of:
 - (a) the amino acid sequence of SEQ ID NO:24;
 - (b) a fragment of the amino acid sequence of SEQ ID NO:24, the fragment comprising eight contiguous amino acids of SEQ ID NO:24; and
- (c) the amino acid sequence encoded by the cDNA insert of clone vrl_1 deposited with the ATCC under accession number 207114; the protein being substantially free from other mammalian proteins.
- 34. An isolated polynucleotide comprising a nucleotide sequence selected from the group consisting of:
 - (a) the nucleotide sequence of SEQ ID NO:25;
 - (b) the nucleotide sequence of SEQ ID NO:25 from nucleotide 13 to nucleotide 513;
 - (c) the nucleotide sequence of the full-length protein coding sequence of clone vc63_1 deposited with the ATCC under accession number 207115;
 - (d) a nucleotide sequence encoding the full-length protein encoded by the cDNA insert of clone vc63_1 deposited with the ATCC under accession number 207115;

(e) a nucleotide sequence encoding a protein comprising the amino acid sequence of SEQ ID NO:26;

- (f) a nucleotide sequence encoding a protein comprising a fragment of the amino acid sequence of SEQ ID NO:26, the fragment comprising eight contiguous amino acids of SEQ ID NO:26;
- (g) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 65 degrees C, or 4X SSC at 42 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(d); and
- (h) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 50 degrees C, or 6X SSC at 40 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(d), and that has a length that is at least 25% of the length of SEQ ID NO:25.
- 35. A protein comprising an amino acid sequence selected from the group consisting of:
 - (a) the amino acid sequence of SEQ ID NO:26;
 - (b) a fragment of the amino acid sequence of SEQ ID NO:26, the fragment comprising eight contiguous amino acids of SEQ ID NO:26; and
- (c) the amino acid sequence encoded by the cDNA insert of clone vc63_1 deposited with the ATCC under accession number 207115; the protein being substantially free from other mammalian proteins.
- 36. An isolated polynucleotide comprising a nucleotide sequence selected from the group consisting of:
 - (a) the nucleotide sequence of SEQ ID NO:27;
 - (b) the nucleotide sequence of SEQ ID NO:27 from nucleotide 79 to nucleotide 345;
 - (c) the nucleotide sequence of SEQ ID NO:27 from nucleotide 130 to nucleotide 345;
 - (d) the nucleotide sequence of the full-length protein coding sequence of clone vb25_1 deposited with the ATCC under accession number PTA-362;

(e) a nucleotide sequence encoding the full-length protein encoded by the cDNA insert of clone vb25_1 deposited with the ATCC under accession number PTA-362;

- (f) the nucleotide sequence of a mature protein coding sequence of clone vb25_1 deposited with the ATCC under accession number PTA-362;
- (g) a nucleotide sequence encoding a mature protein encoded by the cDNA insert of clone vb25_1 deposited with the ATCC under accession number PTA-362;
- (h) a nucleotide sequence encoding a protein comprising the amino acid sequence of SEQ ID NO:28;
- (i) a nucleotide sequence encoding a protein comprising a fragment of the amino acid sequence of SEQ ID NO:28, the fragment comprising eight contiguous amino acids of SEQ ID NO:28;
- (j) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 65 degrees C, or 4X SSC at 42 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g); and
- (k) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 50 degrees C, or 6X SSC at 40 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g), and that has a length that is at least 25% of the length of SEQ ID NO:27.
- 37. A protein comprising an amino acid sequence selected from the group consisting of:
 - (a) the amino acid sequence of SEQ ID NO:28;
 - (b) a fragment of the amino acid sequence of SEQ ID NO:28, the fragment comprising eight contiguous amino acids of SEQ ID NO:28; and
- (c) the amino acid sequence encoded by the cDNA insert of clone vb25_1 deposited with the ATCC under accession number PTA-362; the protein being substantially free from other mammalian proteins.
- 38. An isolated polynucleotide comprising a nucleotide sequence selected from the group consisting of:

- (a) the nucleotide sequence of SEQ ID NO:29;
- (b) the nucleotide sequence of SEQ ID NO:29 from nucleotide 72 to nucleotide 236;
- (c) the nucleotide sequence of SEQ ID NO:29 from nucleotide 150 to nucleotide 236;
- (d) the nucleotide sequence of the full-length protein coding sequence of clone vb27_1 deposited with the ATCC under accession number PTA-362;
- (e) a nucleotide sequence encoding the full-length protein encoded by the cDNA insert of clone vb27_1 deposited with the ATCC under accession number PTA-362;
- (f) the nucleotide sequence of a mature protein coding sequence of clone vb27_1 deposited with the ATCC under accession number PTA-362;
- (g) a nucleotide sequence encoding a mature protein encoded by the cDNA insert of clone vb27_1 deposited with the ATCC under accession number PTA-362;
- (h) a nucleotide sequence encoding a protein comprising the amino acid sequence of SEQ ID NO:30;
- (i) a nucleotide sequence encoding a protein comprising a fragment of the amino acid sequence of SEQ ID NO:30, the fragment comprising eight contiguous amino acids of SEQ ID NO:30;
- (j) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 65 degrees C, or 4X SSC at 42 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g); and
- (k) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 50 degrees C, or 6X SSC at 40 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g), and that has a length that is at least 25% of the length of SEQ ID NO:29.
- 39. A protein comprising an amino acid sequence selected from the group consisting of:
 - (a) the amino acid sequence of SEQ ID NO:30;

(b) a fragment of the amino acid sequence of SEQ ID NO:30, the fragment comprising eight contiguous amino acids of SEQ ID NO:30; and

- (c) the amino acid sequence encoded by the cDNA insert of clone vb27_1 deposited with the ATCC under accession number PTA-362; the protein being substantially free from other mammalian proteins.
- 40. An isolated polynucleotide comprising a nucleotide sequence selected from the group consisting of:
 - (a) the nucleotide sequence of SEQ ID NO:31;
 - (b) the nucleotide sequence of SEQ ID NO:31 from nucleotide 135 to nucleotide 884;
 - (c) the nucleotide sequence of SEQ ID NO:31 from nucleotide 183 to nucleotide 884;
 - (d) the nucleotide sequence of the full-length protein coding sequence of clone vb28_1 deposited with the ATCC under accession number PTA-362;
 - (e) a nucleotide sequence encoding the full-length protein encoded by the cDNA insert of clone vb28_1 deposited with the ATCC under accession number PTA-362;
 - (f) the nucleotide sequence of a mature protein coding sequence of clone vb28_1 deposited with the ATCC under accession number PTA-362;
 - (g) a nucleotide sequence encoding a mature protein encoded by the cDNA insert of clone vb28_1 deposited with the ATCC under accession number PTA-362;
 - (h) a nucleotide sequence encoding a protein comprising the amino acid sequence of SEQ ID NO:32;
 - (i) a nucleotide sequence encoding a protein comprising a fragment of the amino acid sequence of SEQ ID NO:32, the fragment comprising eight contiguous amino acids of SEQ ID NO:32;
 - (j) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 65 degrees C, or 4X SSC at 42 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g); and

(k) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 50 degrees C, or 6X SSC at 40 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g), and that has a length that is at least 25% of the length of SEQ ID NO:31.

- 41. A protein comprising an amino acid sequence selected from the group consisting of:
 - (a) the amino acid sequence of SEQ ID NO:32;
 - (b) a fragment of the amino acid sequence of SEQ ID NO:32, the fragment comprising eight contiguous amino acids of SEQ ID NO:32; and
- (c) the amino acid sequence encoded by the cDNA insert of clone vb28_1 deposited with the ATCC under accession number PTA-362; the protein being substantially free from other mammalian proteins.
- 42. An isolated polynucleotide comprising a nucleotide sequence selected from the group consisting of:
 - (a) the nucleotide sequence of SEQ ID NO:33;
 - (b) the nucleotide sequence of SEQ ID NO:33 from nucleotide 42 to nucleotide 206;
 - (c) the nucleotide sequence of SEQ ID NO:33 from nucleotide 111 to nucleotide 206;
 - (d) the nucleotide sequence of the full-length protein coding sequence of clone vb29_1 deposited with the ATCC under accession number PTA-362;
 - (e) a nucleotide sequence encoding the full-length protein encoded by the cDNA insert of clone vb29_1 deposited with the ATCC under accession number PTA-362;
 - (f) the nucleotide sequence of a mature protein coding sequence of clone vb29_1 deposited with the ATCC under accession number PTA-362;
 - (g) a nucleotide sequence encoding a mature protein encoded by the cDNA insert of clone vb29_1 deposited with the ATCC under accession number PTA-362;

(h) a nucleotide sequence encoding a protein comprising the amino acid sequence of SEQ ID NO:34;

- (i) a nucleotide sequence encoding a protein comprising a fragment of the amino acid sequence of SEQ ID NO:34, the fragment comprising eight contiguous amino acids of SEQ ID NO:34;
- (j) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 65 degrees C, or 4X SSC at 42 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g); and
- (k) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 50 degrees C, or 6X SSC at 40 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g), and that has a length that is at least 25% of the length of SEQ ID NO:33.
- 43. A protein comprising an amino acid sequence selected from the group consisting of:
 - (a) the amino acid sequence of SEQ ID NO:34;
 - (b) a fragment of the amino acid sequence of SEQ ID NO:34, the fragment comprising eight contiguous amino acids of SEQ ID NO:34; and
- (c) the amino acid sequence encoded by the cDNA insert of clone vb29_1 deposited with the ATCC under accession number PTA-362; the protein being substantially free from other mammalian proteins.
- 44. An isolated polynucleotide comprising a nucleotide sequence selected from the group consisting of:
 - (a) a polynucleotide comprising the nucleotide sequence of SEQ ID NO:35;
 - (b) a polynucleotide comprising the nucleotide sequence of SEQ ID NO:35 from nucleotide 17 to nucleotide 253;
 - (c) a polynucleotide comprising the nucleotide sequence of SEQ ID NO:35 from nucleotide 98 to nucleotide 253;

(d) a polynucleotide comprising the nucleotide sequence of the full-length protein coding sequence of clone vb30_1 deposited with the ATCC under accession number PTA-362;

- (e) a polynucleotide encoding the full-length protein encoded by the cDNA insert of clone vb30_1 deposited with the ATCC under accession number PTA-362;
- (f) a polynucleotide comprising the nucleotide sequence of a mature protein coding sequence of clone vb30_1 deposited with the ATCC under accession number PTA-362;
- (g) a polynucleotide encoding a mature protein encoded by the cDNA insert of clone vb30_1 deposited with the ATCC under accession number PTA-362;
- (h) a polynucleotide encoding a protein comprising the amino acid sequence of SEQ ID NO:36;
- (i) a polynucleotide encoding a protein comprising a fragment of the amino acid sequence of SEQ ID NO:36 having biological activity, the fragment comprising eight contiguous amino acids of SEQ ID NO:36;
- (j) a polynucleotide which is an allelic variant of a polynucleotide of(a)-(g) above;
- (k) a polynucleotide which encodes a species homologue of the protein of (h) or (i) above;
- (l) a polynucleotide that hybridizes under stringent conditions to any one of the polynucleotides specified in (a)-(i); and
- (m) a polynucleotide that hybridizes under stringent conditions to any one of the polynucleotides specified in (a)-(i) and that has a length that is at least 25% of the length of SEQ ID NO:35.
- 45. A protein comprising an amino acid sequence selected from the group consisting of:
 - (a) the amino acid sequence of SEQ ID NO:36;
 - (b) a fragment of the amino acid sequence of SEQ ID NO:36, the fragment comprising eight contiguous amino acids of SEQ ID NO:36; and

(c) the amino acid sequence encoded by the cDNA insert of clone vb30_1 deposited with the ATCC under accession number PTA-362; the protein being substantially free from other mammalian proteins.

- 46. An isolated polynucleotide comprising a nucleotide sequence selected from the group consisting of:
 - (a) the nucleotide sequence of SEQ ID NO:37;
 - (b) the nucleotide sequence of SEQ ID NO:37 from nucleotide 68 to nucleotide 424;
 - (c) the nucleotide sequence of the full-length protein coding sequence of clone vc67_1 deposited with the ATCC under accession number PTA-362;
 - (d) a nucleotide sequence encoding the full-length protein encoded by the cDNA insert of clone vc67_1 deposited with the ATCC under accession number PTA-362;
 - (e) a nucleotide sequence encoding a protein comprising the amino acid sequence of SEQ ID NO:38;
 - (f) a nucleotide sequence encoding a protein comprising a fragment of the amino acid sequence of SEQ ID NO:38, the fragment comprising eight contiguous amino acids of SEQ ID NO:38;
 - (g) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 65 degrees C, or 4X SSC at 42 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(d); and
 - (h) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 50 degrees C, or 6X SSC at 40 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(d), and that has a length that is at least 25% of the length of SEQ ID NO:37.
- 47. A protein comprising an amino acid sequence selected from the group consisting of:
 - (a) the amino acid sequence of SEQ ID NO:38;
 - (b) a fragment of the amino acid sequence of SEQ ID NO:38, the fragment comprising eight contiguous amino acids of SEQ ID NO:38; and

(c) the amino acid sequence encoded by the cDNA insert of clone vc67_1 deposited with the ATCC under accession number PTA-362; the protein being substantially free from other mammalian proteins.

- 48. An isolated polynucleotide comprising a nucleotide sequence selected from the group consisting of:
 - (a) the nucleotide sequence of SEQ ID NO:39;
 - (b) the nucleotide sequence of SEQ ID NO:39 from nucleotide 103 to nucleotide 261;
 - (c) the nucleotide sequence of SEQ ID NO:39 from nucleotide 154 to nucleotide 261;
 - (d) the nucleotide sequence of the full-length protein coding sequence of clone vf4_1 deposited with the ATCC under accession number PTA-362;
 - (e) a nucleotide sequence encoding the full-length protein encoded by the cDNA insert of clone vf4_1 deposited with the ATCC under accession number PTA-362;
 - (f) the nucleotide sequence of a mature protein coding sequence of clone vf4_1 deposited with the ATCC under accession number PTA-362;
 - (g) a nucleotide sequence encoding a mature protein encoded by the cDNA insert of clone vf4_1 deposited with the ATCC under accession number PTA-362;
 - (h) a nucleotide sequence encoding a protein comprising the amino acid sequence of SEQ ID NO:40;
 - (i) a nucleotide sequence encoding a protein comprising a fragment of the amino acid sequence of SEQ ID NO:40, the fragment comprising eight contiguous amino acids of SEQ ID NO:40;
 - (j) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 65 degrees C, or 4X SSC at 42 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g); and
 - (k) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 50 degrees C, or 6X SSC at 40 degrees

C with 50% formamide, to any one of the polynucleotides specified by (a)-(g), and that has a length that is at least 25% of the length of SEQ ID NO:39.

- 49. A protein comprising an amino acid sequence selected from the group consisting of:
 - (a) the amino acid sequence of SEQ ID NO:40;
 - (b) a fragment of the amino acid sequence of SEQ ID NO:40, the fragment comprising eight contiguous amino acids of SEQ ID NO:40; and
- (c) the amino acid sequence encoded by the cDNA insert of clone vf4_1 deposited with the ATCC under accession number PTA-362; the protein being substantially free from other mammalian proteins.
- 50. An isolated polynucleotide comprising a nucleotide sequence selected from the group consisting of:
 - (a) the nucleotide sequence of SEQ ID NO:41;
 - (b) the nucleotide sequence of SEQ ID NO:41 from nucleotide 1575 to nucleotide 3038;
 - (c) the nucleotide sequence of SEQ ID NO:41 from nucleotide 1650 to nucleotide 3038;
 - (d) the nucleotide sequence of the full-length protein coding sequence of clone vg3_1 deposited with the ATCC under accession number PTA-362;
 - (e) a nucleotide sequence encoding the full-length protein encoded by the cDNA insert of clone vg3_1 deposited with the ATCC under accession number PTA-362;
 - (f) the nucleotide sequence of a mature protein coding sequence of clone vg3_1 deposited with the ATCC under accession number PTA-362;
 - (g) a nucleotide sequence encoding a mature protein encoded by the cDNA insert of clone vg3_1 deposited with the ATCC under accession number PTA-362;
 - (h) a nucleotide sequence encoding a protein comprising the amino acid sequence of SEQ ID NO:42;

(i) a nucleotide sequence encoding a protein comprising a fragment of the amino acid sequence of SEQ ID NO:42, the fragment comprising eight contiguous amino acids of SEQ ID NO:42;

- (j) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 65 degrees C, or 4X SSC at 42 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g); and
- (k) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 50 degrees C, or 6X SSC at 40 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g), and that has a length that is at least 25% of the length of SEQ ID NO:41.
- 51. A protein comprising an amino acid sequence selected from the group consisting of:
 - (a) the amino acid sequence of SEQ ID NO:42;
 - (b) a fragment of the amino acid sequence of SEQ ID NO:42, the fragment comprising eight contiguous amino acids of SEQ ID NO:42; and
- (c) the amino acid sequence encoded by the cDNA insert of clone vg3_1 deposited with the ATCC under accession number PTA-362; the protein being substantially free from other mammalian proteins.
- 52. An isolated polynucleotide comprising a nucleotide sequence selected from the group consisting of:
 - (a) the nucleotide sequence of SEQ ID NO:43;
 - (b) the nucleotide sequence of SEQ ID NO:43 from nucleotide 2112 to nucleotide 2363;
 - (c) the nucleotide sequence of the full-length protein coding sequence of clone vo2_1 deposited with the ATCC under accession number PTA-362;
 - (d) a nucleotide sequence encoding the full-length protein encoded by the cDNA insert of clone vo2_1 deposited with the ATCC under accession number PTA-362;
 - (e) a nucleotide sequence encoding a protein comprising the amino acid sequence of SEQ ID NO:44;

(f) a nucleotide sequence encoding a protein comprising a fragment of the amino acid sequence of SEQ ID NO:44, the fragment comprising eight contiguous amino acids of SEQ ID NO:44;

- (g) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 65 degrees C, or 4X SSC at 42 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(d); and
- (h) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 50 degrees C, or 6X SSC at 40 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(d), and that has a length that is at least 25% of the length of SEQ ID NO:43.
- 53. A protein comprising an amino acid sequence selected from the group consisting of:
 - (a) the amino acid sequence of SEQ ID NO:44;
 - (b) a fragment of the amino acid sequence of SEQ ID NO:44, the fragment comprising eight contiguous amino acids of SEQ ID NO:44; and
- (c) the amino acid sequence encoded by the cDNA insert of clone vo2_1 deposited with the ATCC under accession number PTA-362; the protein being substantially free from other mammalian proteins.
- 54. An isolated polynucleotide comprising a nucleotide sequence selected from the group consisting of:
 - (a) the nucleotide sequence of SEQ ID NO:45;
 - (b) the nucleotide sequence of SEQ ID NO:45 from nucleotide 36 to nucleotide 707;
 - (c) the nucleotide sequence of SEQ ID NO:45 from nucleotide 393 to nucleotide 707;
 - (d) the nucleotide sequence of the full-length protein coding sequence of clone vo3_1 deposited with the ATCC under accession number PTA-362;
 - (e) a nucleotide sequence encoding the full-length protein encoded by the cDNA insert of clone vo3_1 deposited with the ATCC under accession number PTA-362;

(f) the nucleotide sequence of a mature protein coding sequence of clone vo3_1 deposited with the ATCC under accession number PTA-362;

- (g) a nucleotide sequence encoding a mature protein encoded by the cDNA insert of clone vo3_1 deposited with the ATCC under accession number PTA-362;
- (h) a nucleotide sequence encoding a protein comprising the amino acid sequence of SEQ ID NO:46;
- (i) a nucleotide sequence encoding a protein comprising a fragment of the amino acid sequence of SEQ ID NO:46, the fragment comprising eight contiguous amino acids of SEQ ID NO:46;
- (j) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 65 degrees C, or 4X SSC at 42 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g); and
- (k) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 50 degrees C, or 6X SSC at 40 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g), and that has a length that is at least 25% of the length of SEQ ID NO:45.
- 55. A protein comprising an amino acid sequence selected from the group consisting of:
 - (a) the amino acid sequence of SEQ ID NO:46;
 - (b) a fragment of the amino acid sequence of SEQ ID NO:46, the fragment comprising eight contiguous amino acids of SEQ ID NO:46; and
- (c) the amino acid sequence encoded by the cDNA insert of clone vo3_1 deposited with the ATCC under accession number PTA-362; the protein being substantially free from other mammalian proteins.
- 56. An isolated polynucleotide comprising a nucleotide sequence selected from the group consisting of:
 - (a) the nucleotide sequence of SEQ ID NO:47;
 - (b) the nucleotide sequence of SEQ ID NO:47 from nucleotide 74 to nucleotide 295;

(c) the nucleotide sequence of SEQ ID NO:47 from nucleotide 134 to nucleotide 295;

- (d) the nucleotide sequence of the full-length protein coding sequence of clone vo5_1 deposited with the ATCC under accession number PTA-362;
- (e) a nucleotide sequence encoding the full-length protein encoded by the cDNA insert of clone vo5_1 deposited with the ATCC under accession number PTA-362;
- (f) the nucleotide sequence of a mature protein coding sequence of clone vo5_1 deposited with the ATCC under accession number PTA-362;
- (g) a nucleotide sequence encoding a mature protein encoded by the cDNA insert of clone vo5_1 deposited with the ATCC under accession number PTA-362;
- (h) a nucleotide sequence encoding a protein comprising the amino acid sequence of SEQ ID NO:48;
- (i) a nucleotide sequence encoding a protein comprising a fragment of the amino acid sequence of SEQ ID NO:48, the fragment comprising eight contiguous amino acids of SEQ ID NO:48;
- (j) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 65 degrees C, or 4X SSC at 42 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g); and
- (k) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 50 degrees C, or 6X SSC at 40 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g), and that has a length that is at least 25% of the length of SEQ ID NO:47.
- 57. A protein comprising an amino acid sequence selected from the group consisting of:
 - (a) the amino acid sequence of SEQ ID NO:48;
 - (b) a fragment of the amino acid sequence of SEQ ID NO:48, the fragment comprising eight contiguous amino acids of SEQ ID NO:48; and
 - (c) the amino acid sequence encoded by the cDNA insert of clone vo5_1 deposited with the ATCC under accession number PTA-362;

the protein being substantially free from other mammalian proteins.

58. An isolated polynucleotide comprising a nucleotide sequence selected from the group consisting of:

- (a) the nucleotide sequence of SEQ ID NO:49;
- (b) the nucleotide sequence of SEQ ID NO:49 from nucleotide 45 to nucleotide 383;
- (c) the nucleotide sequence of SEQ ID NO:49 from nucleotide 312 to nucleotide 383;
- (d) the nucleotide sequence of the full-length protein coding sequence of clone vo6_1 deposited with the ATCC under accession number PTA-362;
- (e) a nucleotide sequence encoding the full-length protein encoded by the cDNA insert of clone vo6_1 deposited with the ATCC under accession number PTA-362;
- (f) the nucleotide sequence of a mature protein coding sequence of clone vo6_1 deposited with the ATCC under accession number PTA-362;
- (g) a nucleotide sequence encoding a mature protein encoded by the cDNA insert of clone vo6_1 deposited with the ATCC under accession number PTA-362;
- (h) a nucleotide sequence encoding a protein comprising the amino acid sequence of SEQ ID NO:50;
- (i) a nucleotide sequence encoding a protein comprising a fragment of the amino acid sequence of SEQ ID NO:50, the fragment comprising eight contiguous amino acids of SEQ ID NO:50;
- (j) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 65 degrees C, or 4X SSC at 42 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g); and
- (k) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 50 degrees C, or 6X SSC at 40 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g), and that has a length that is at least 25% of the length of SEQ ID NO:49.

59. A protein comprising an amino acid sequence selected from the group consisting of:

- (a) the amino acid sequence of SEQ ID NO:50;
- (b) a fragment of the amino acid sequence of SEQ ID NO:50, the fragment comprising eight contiguous amino acids of SEQ ID NO:50; and
- (c) the amino acid sequence encoded by the cDNA insert of clone vo6_1 deposited with the ATCC under accession number PTA-362; the protein being substantially free from other mammalian proteins.
- 60. An isolated polynucleotide comprising a nucleotide sequence selected from the group consisting of:
 - (a) the nucleotide sequence of SEQ ID NO:51;
 - (b) the nucleotide sequence of SEQ ID NO:51 from nucleotide 186 to nucleotide 1739;
 - (c) the nucleotide sequence of SEQ ID NO:51 from nucleotide 288 to nucleotide 1739;
 - (d) the nucleotide sequence of the full-length protein coding sequence of clone vo9_1 deposited with the ATCC under accession number PTA-362;
 - (e) a nucleotide sequence encoding the full-length protein encoded by the cDNA insert of clone vo9_1 deposited with the ATCC under accession number PTA-362:
 - (f) the nucleotide sequence of a mature protein coding sequence of clone vo9_1 deposited with the ATCC under accession number PTA-362;
 - (g) a nucleotide sequence encoding a mature protein encoded by the cDNA insert of clone vo9_1 deposited with the ATCC under accession number PTA-362;
 - (h) a nucleotide sequence encoding a protein comprising the amino acid sequence of SEQ ID NO:52;
 - (i) a nucleotide sequence encoding a protein comprising a fragment of the amino acid sequence of SEQ ID NO:52, the fragment comprising eight contiguous amino acids of SEQ ID NO:52;

(j) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 65 degrees C, or 4X SSC at 42 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g); and

- (k) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 50 degrees C, or 6X SSC at 40 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g), and that has a length that is at least 25% of the length of SEQ ID NO:51.
- 61. A protein comprising an amino acid sequence selected from the group consisting of:
 - (a) the amino acid sequence of SEQ ID NO:52;
 - (b) a fragment of the amino acid sequence of SEQ ID NO:52, the fragment comprising eight contiguous amino acids of SEQ ID NO:52; and
- (c) the amino acid sequence encoded by the cDNA insert of clone vo9_1 deposited with the ATCC under accession number PTA-362; the protein being substantially free from other mammalian proteins.
- 62. An isolated polynucleotide comprising a nucleotide sequence selected from the group consisting of:
 - (a) the nucleotide sequence of SEQ ID NO:53;
 - (b) the nucleotide sequence of SEQ ID NO:53 from nucleotide 440 to nucleotide 835;
 - (c) the nucleotide sequence of SEQ ID NO:53 from nucleotide 632 to nucleotide 835;
 - (d) the nucleotide sequence of the full-length protein coding sequence of clone vol1_1 deposited with the ATCC under accession number PTA-366;
 - (e) a nucleotide sequence encoding the full-length protein encoded by the cDNA insert of clone vol1_1 deposited with the ATCC under accession number PTA-366;
 - (f) the nucleotide sequence of a mature protein coding sequence of clone vol1_1 deposited with the ATCC under accession number PTA-366;

(g) a nucleotide sequence encoding a mature protein encoded by the cDNA insert of clone vol1_1 deposited with the ATCC under accession number PTA-366;

- (h) a nucleotide sequence encoding a protein comprising the amino acid sequence of SEQ ID NO:54;
- (i) a nucleotide sequence encoding a protein comprising a fragment of the amino acid sequence of SEQ ID NO:54, the fragment comprising eight contiguous amino acids of SEQ ID NO:54;
- (j) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 65 degrees C, or 4X SSC at 42 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g); and
- (k) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 50 degrees C, or 6X SSC at 40 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g), and that has a length that is at least 25% of the length of SEQ ID NO:53.
- 63. A protein comprising an amino acid sequence selected from the group consisting of:
 - (a) the amino acid sequence of SEQ ID NO:54;
 - (b) a fragment of the amino acid sequence of SEQ ID NO:54, the fragment comprising eight contiguous amino acids of SEQ ID NO:54; and
- (c) the amino acid sequence encoded by the cDNA insert of clone vol1_1 deposited with the ATCC under accession number PTA-366; the protein being substantially free from other mammalian proteins.
- 64. An isolated polynucleotide comprising a nucleotide sequence selected from the group consisting of:
 - (a) the nucleotide sequence of SEQ ID NO:55;
 - (b) the nucleotide sequence of SEQ ID NO:55 from nucleotide 72 to nucleotide 329;
 - (c) the nucleotide sequence of SEQ ID NO:55 from nucleotide 120 to nucleotide 329;

(d) the nucleotide sequence of the full-length protein coding sequence of clone vo12_1 deposited with the ATCC under accession number PTA-366;

- (e) a nucleotide sequence encoding the full-length protein encoded by the cDNA insert of clone vo12_1 deposited with the ATCC under accession number PTA-366;
- (f) the nucleotide sequence of a mature protein coding sequence of clone vo12_1 deposited with the ATCC under accession number PTA-366;
- (g) a nucleotide sequence encoding a mature protein encoded by the cDNA insert of clone vo12_1 deposited with the ATCC under accession number PTA-366;
- (h) a nucleotide sequence encoding a protein comprising the amino acid sequence of SEQ ID NO:56;
- (i) a nucleotide sequence encoding a protein comprising a fragment of the amino acid sequence of SEQ ID NO:56, the fragment comprising eight contiguous amino acids of SEQ ID NO:56;
- (j) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 65 degrees C, or 4X SSC at 42 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g); and
- (k) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 50 degrees C, or 6X SSC at 40 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g), and that has a length that is at least 25% of the length of SEQ ID NO:55.
- 65. A protein comprising an amino acid sequence selected from the group consisting of:
 - (a) the amino acid sequence of SEQ ID NO:56;
 - (b) a fragment of the amino acid sequence of SEQ ID NO:56, the fragment comprising eight contiguous amino acids of SEQ ID NO:56; and
- (c) the amino acid sequence encoded by the cDNA insert of clone vo12_1 deposited with the ATCC under accession number PTA-366; the protein being substantially free from other mammalian proteins.

66. An isolated polynucleotide comprising a nucleotide sequence selected from the group consisting of:

- (a) the nucleotide sequence of SEQ ID NO:57;
- (b) the nucleotide sequence of SEQ ID NO:57 from nucleotide 227 to nucleotide 439;
- (c) the nucleotide sequence of SEQ ID NO:57 from nucleotide 287 to nucleotide 439;
- (d) the nucleotide sequence of the full-length protein coding sequence of clone vo13_1 deposited with the ATCC under accession number PTA-366;
- (e) a nucleotide sequence encoding the full-length protein encoded by the cDNA insert of clone vo13_1 deposited with the ATCC under accession number PTA-366;
- (f) the nucleotide sequence of a mature protein coding sequence of clone vol3_1 deposited with the ATCC under accession number PTA-366;
- (g) a nucleotide sequence encoding a mature protein encoded by the cDNA insert of clone vol3_1 deposited with the ATCC under accession number PTA-366;
- (h) a nucleotide sequence encoding a protein comprising the amino acid sequence of SEQ ID NO:58;
- (i) a nucleotide sequence encoding a protein comprising a fragment of the amino acid sequence of SEQ ID NO:58, the fragment comprising eight contiguous amino acids of SEQ ID NO:58;
- (j) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 65 degrees C, or 4X SSC at 42 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g); and
- (k) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 50 degrees C, or 6X SSC at 40 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g), and that has a length that is at least 25% of the length of SEQ ID NO:57.
- 67. A protein comprising an amino acid sequence selected from the group consisting of:

(a) the amino acid sequence of SEQ ID NO:58;

- (b) a fragment of the amino acid sequence of SEQ ID NO:58, the fragment comprising eight contiguous amino acids of SEQ ID NO:58; and
- (c) the amino acid sequence encoded by the cDNA insert of clone vo13_1 deposited with the ATCC under accession number PTA-366; the protein being substantially free from other mammalian proteins.
- 68. An isolated polynucleotide comprising a nucleotide sequence selected from the group consisting of:
 - (a) the nucleotide sequence of SEQ ID NO:59;
 - (b) the nucleotide sequence of SEQ ID NO:59 from nucleotide 96 to nucleotide 341;
 - (c) the nucleotide sequence of SEQ ID NO:59 from nucleotide 174 to nucleotide 341;
 - (d) the nucleotide sequence of the full-length protein coding sequence of clone vo14_1 deposited with the ATCC under accession number PTA-366;
 - (e) a nucleotide sequence encoding the full-length protein encoded by the cDNA insert of clone vo14_1 deposited with the ATCC under accession number PTA-366;
 - (f) the nucleotide sequence of a mature protein coding sequence of clone vo14_1 deposited with the ATCC under accession number PTA-366;
 - (g) a nucleotide sequence encoding a mature protein encoded by the cDNA insert of clone vo14_1 deposited with the ATCC under accession number PTA-366;
 - (h) a nucleotide sequence encoding a protein comprising the amino acid sequence of SEQ ID NO:60;
 - (i) a nucleotide sequence encoding a protein comprising a fragment of the amino acid sequence of SEQ ID NO:60, the fragment comprising eight contiguous amino acids of SEQ ID NO:60;
 - (j) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 65 degrees C, or 4X SSC at 42 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g); and

(k) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 50 degrees C, or 6X SSC at 40 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g), and that has a length that is at least 25% of the length of SEQ ID NO:59.

- 69. A protein comprising an amino acid sequence selected from the group consisting of:
 - (a) the amino acid sequence of SEQ ID NO:60;
 - (b) a fragment of the amino acid sequence of SEQ ID NO:60, the fragment comprising eight contiguous amino acids of SEQ ID NO:60; and
- (c) the amino acid sequence encoded by the cDNA insert of clone vo14_1 deposited with the ATCC under accession number PTA-366; the protein being substantially free from other mammalian proteins.
- 70. An isolated polynucleotide comprising a nucleotide sequence selected from the group consisting of:
 - (a) the nucleotide sequence of SEQ ID NO:61;
 - (b) the nucleotide sequence of SEQ ID NO:61 from nucleotide 90 to nucleotide 599;
 - (c) the nucleotide sequence of SEQ ID NO:61 from nucleotide 165 to nucleotide 599;
 - (d) the nucleotide sequence of the full-length protein coding sequence of clone vo15_1 deposited with the ATCC under accession number PTA-366;
 - (e) a nucleotide sequence encoding the full-length protein encoded by the cDNA insert of clone vo15_1 deposited with the ATCC under accession number PTA-366;
 - (f) the nucleotide sequence of a mature protein coding sequence of clone vo15_1 deposited with the ATCC under accession number PTA-366;
 - (g) a nucleotide sequence encoding a mature protein encoded by the cDNA insert of clone vo15_1 deposited with the ATCC under accession number PTA-366;

(h) a nucleotide sequence encoding a protein comprising the amino acid sequence of SEQ ID NO:62;

- (i) a nucleotide sequence encoding a protein comprising a fragment of the amino acid sequence of SEQ ID NO:62, the fragment comprising eight contiguous amino acids of SEQ ID NO:62;
- (j) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 65 degrees C, or 4X SSC at 42 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g); and
- (k) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 50 degrees C, or 6X SSC at 40 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g), and that has a length that is at least 25% of the length of SEQ ID NO:61.
- 71. A protein comprising an amino acid sequence selected from the group consisting of:
 - (a) the amino acid sequence of SEQ ID NO:62;
 - (b) a fragment of the amino acid sequence of SEQ ID NO:62, the fragment comprising eight contiguous amino acids of SEQ ID NO:62; and
- (c) the amino acid sequence encoded by the cDNA insert of clone vo15_1 deposited with the ATCC under accession number PTA-366; the protein being substantially free from other mammalian proteins.
- 72. An isolated polynucleotide comprising a nucleotide sequence selected from the group consisting of:
 - (a) the nucleotide sequence of SEQ ID NO:63;
 - (b) the nucleotide sequence of SEQ ID NO:63 from nucleotide 209 to nucleotide 451;
 - (c) the nucleotide sequence of SEQ ID NO:63 from nucleotide 398 to nucleotide 451;
 - (d) the nucleotide sequence of the full-length protein coding sequence of clone vo16_1 deposited with the ATCC under accession number PTA-366;

(e) a nucleotide sequence encoding the full-length protein encoded by the cDNA insert of clone vo16_1 deposited with the ATCC under accession number PTA-366;

- (f) the nucleotide sequence of a mature protein coding sequence of clone vo16_1 deposited with the ATCC under accession number PTA-366;
- (g) a nucleotide sequence encoding a mature protein encoded by the cDNA insert of clone vo16_1 deposited with the ATCC under accession number PTA-366;
- (h) a nucleotide sequence encoding a protein comprising the amino acid sequence of SEQ ID NO:64;
- (i) a nucleotide sequence encoding a protein comprising a fragment of the amino acid sequence of SEQ ID NO:64, the fragment comprising eight contiguous amino acids of SEQ ID NO:64;
- (j) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 65 degrees C, or 4X SSC at 42 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g); and
- (k) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 50 degrees C, or 6X SSC at 40 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g), and that has a length that is at least 25% of the length of SEQ ID NO:63.
- 73. A protein comprising an amino acid sequence selected from the group consisting of:
 - (a) the amino acid sequence of SEQ ID NO:64;
 - (b) a fragment of the amino acid sequence of SEQ ID NO:64, the fragment comprising eight contiguous amino acids of SEQ ID NO:64; and
- (c) the amino acid sequence encoded by the cDNA insert of clone vo16_1 deposited with the ATCC under accession number PTA-366; the protein being substantially free from other mammalian proteins.
- 74. An isolated polynucleotide comprising a nucleotide sequence selected from the group consisting of:

- (a) the nucleotide sequence of SEQ ID NO:65;
- (b) the nucleotide sequence of SEQ ID NO:65 from nucleotide 31 to nucleotide 231;
- (c) the nucleotide sequence of SEQ ID NO:65 from nucleotide 97 to nucleotide 231;
- (d) the nucleotide sequence of the full-length protein coding sequence of clone vo18_1 deposited with the ATCC under accession number PTA-366;
- (e) a nucleotide sequence encoding the full-length protein encoded by the cDNA insert of clone vo18_1 deposited with the ATCC under accession number PTA-366;
- (f) the nucleotide sequence of a mature protein coding sequence of clone vo18_1 deposited with the ATCC under accession number PTA-366;
- (g) a nucleotide sequence encoding a mature protein encoded by the cDNA insert of clone vo18_1 deposited with the ATCC under accession number, PTA-366;
- (h) a nucleotide sequence encoding a protein comprising the amino acid sequence of SEQ ID NO:66;
- (i) a nucleotide sequence encoding a protein comprising a fragment of the amino acid sequence of SEQ ID NO:66, the fragment comprising eight contiguous amino acids of SEQ ID NO:66;
- (j) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 65 degrees C, or 4X SSC at 42 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g); and
- (k) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 50 degrees C, or 6X SSC at 40 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g), and that has a length that is at least 25% of the length of SEQ ID NO:65.
- 75. A protein comprising an amino acid sequence selected from the group consisting of:
 - (a) the amino acid sequence of SEQ ID NO:66;

(b) a fragment of the amino acid sequence of SEQ ID NO:66, the fragment comprising eight contiguous amino acids of SEQ ID NO:66; and

- (c) the amino acid sequence encoded by the cDNA insert of clone vo18_1 deposited with the ATCC under accession number PTA-366; the protein being substantially free from other mammalian proteins.
- 76. An isolated polynucleotide comprising a nucleotide sequence selected from the group consisting of:
 - (a) the nucleotide sequence of SEQ ID NO:67;
 - (b) the nucleotide sequence of SEQ ID NO:67 from nucleotide 23 to nucleotide 736;
 - (c) the nucleotide sequence of SEQ ID NO:67 from nucleotide 83 to nucleotide 736;
 - (d) the nucleotide sequence of the full-length protein coding sequence of clone vo19_1 deposited with the ATCC under accession number PTA-366;
 - (e) a nucleotide sequence encoding the full-length protein encoded by the cDNA insert of clone vo19_1 deposited with the ATCC under accession number PTA-366;
 - (f) the nucleotide sequence of a mature protein coding sequence of clone vo19_1 deposited with the ATCC under accession number PTA-366;
 - (g) a nucleotide sequence encoding a mature protein encoded by the cDNA insert of clone vo19_1 deposited with the ATCC under accession number PTA-366;
 - (h) a nucleotide sequence encoding a protein comprising the amino acid sequence of SEQ ID NO:68;
 - (i) a nucleotide sequence encoding a protein comprising a fragment of the amino acid sequence of SEQ ID NO:68, the fragment comprising eight contiguous amino acids of SEQ ID NO:68;
 - (j) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 65 degrees C, or 4X SSC at 42 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g); and

(k) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 50 degrees C, or 6X SSC at 40 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g), and that has a length that is at least 25% of the length of SEQ ID NO:67.

- 77. A protein comprising an amino acid sequence selected from the group consisting of:
 - (a) the amino acid sequence of SEQ ID NO:68;
 - (b) a fragment of the amino acid sequence of SEQ ID NO:68, the fragment comprising eight contiguous amino acids of SEQ ID NO:68; and
- (c) the amino acid sequence encoded by the cDNA insert of clone vo19_1 deposited with the ATCC under accession number PTA-366; the protein being substantially free from other mammalian proteins.
- 78. An isolated polynucleotide comprising a nucleotide sequence selected from the group consisting of:
 - (a) the nucleotide sequence of SEQ ID NO:69;
 - (b) the nucleotide sequence of SEQ ID NO:69 from nucleotide 104 to nucleotide 1399;
 - (c) the nucleotide sequence of SEQ ID NO:69 from nucleotide 158 to nucleotide 1399;
 - (d) the nucleotide sequence of the full-length protein coding sequence of clone vo22_1 deposited with the ATCC under accession number PTA-366;
 - (e) a nucleotide sequence encoding the full-length protein encoded by the cDNA insert of clone vo22_1 deposited with the ATCC under accession number PTA-366;
 - (f) the nucleotide sequence of a mature protein coding sequence of clone vo22_1 deposited with the ATCC under accession number PTA-366;
 - (g) a nucleotide sequence encoding a mature protein encoded by the cDNA insert of clone vo22_1 deposited with the ATCC under accession number PTA-366;

(h) a nucleotide sequence encoding a protein comprising the amino acid sequence of SEQ ID NO:70;

- (i) a nucleotide sequence encoding a protein comprising a fragment of the amino acid sequence of SEQ ID NO:70, the fragment comprising eight contiguous amino acids of SEQ ID NO:70;
- (j) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 65 degrees C, or 4X SSC at 42 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g); and
- (k) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 50 degrees C, or 6X SSC at 40 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g), and that has a length that is at least 25% of the length of SEQ ID NO:69.
- 79. A protein comprising an amino acid sequence selected from the group consisting of:
 - (a) the amino acid sequence of SEQ ID NO:70;
 - (b) a fragment of the amino acid sequence of SEQ ID NO:70, the fragment comprising eight contiguous amino acids of SEQ ID NO:70; and
- (c) the amino acid sequence encoded by the cDNA insert of clone vo22_1 deposited with the ATCC under accession number PTA-366; the protein being substantially free from other mammalian proteins.
- 80. An isolated polynucleotide comprising a nucleotide sequence selected from the group consisting of:
 - (a) the nucleotide sequence of SEQ ID NO:71;
 - (b) the nucleotide sequence of SEQ ID NO:71 from nucleotide 174 to nucleotide 1595;
 - (c) the nucleotide sequence of the full-length protein coding sequence of clone vo23_1 deposited with the ATCC under accession number PTA-366;
 - (d) a nucleotide sequence encoding the full-length protein encoded by the cDNA insert of clone vo23_1 deposited with the ATCC under accession number PTA-366;

(e) a nucleotide sequence encoding a protein comprising the amino acid sequence of SEQ ID NO:72;

- (f) a nucleotide sequence encoding a protein comprising a fragment of the amino acid sequence of SEQ ID NO:72, the fragment comprising eight contiguous amino acids of SEQ ID NO:72;
- (g) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 65 degrees C, or 4X SSC at 42 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(d); and
- (h) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 50 degrees C, or 6X SSC at 40 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(d), and that has a length that is at least 25% of the length of SEQ ID NO:71.
- 81. A protein comprising an amino acid sequence selected from the group consisting of:
 - (a) the amino acid sequence of SEQ ID NO:72;
 - (b) a fragment of the amino acid sequence of SEQ ID NO:72, the fragment comprising eight contiguous amino acids of SEQ ID NO:72; and
- (c) the amino acid sequence encoded by the cDNA insert of clone vo23_1 deposited with the ATCC under accession number PTA-366; the protein being substantially free from other mammalian proteins.
- 82. An isolated polynucleotide comprising a nucleotide sequence selected from the group consisting of:
 - (a) the nucleotide sequence of SEQ ID NO:73;
 - (b) the nucleotide sequence of SEQ ID NO:73 from nucleotide 129 to nucleotide 311;
 - (c) the nucleotide sequence of SEQ ID NO:73 from nucleotide 195 to nucleotide 311;
 - (d) the nucleotide sequence of the full-length protein coding sequence of clone vo24_1 deposited with the ATCC under accession number PTA-366;

PCT/US09/07285

- (e) a nucleotide sequence encoding the full-length protein encoded by the cDNA insert of clone vo24_1 deposited with the ATCC under accession number PTA-366;
- (f) the nucleotide sequence of a mature protein coding sequence of clone vo24_1 deposited with the ATCC under accession number PTA-366;
- (g) a nucleotide sequence encoding a mature protein encoded by the cDNA insert of clone vo24_1 deposited with the ATCC under accession number PTA-366;
- (h) a nucleotide sequence encoding a protein comprising the amino acid sequence of SEQ ID NO:74;
- (i) a nucleotide sequence encoding a protein comprising a fragment of the amino acid sequence of SEQ ID NO:74, the fragment comprising eight contiguous amino acids of SEQ ID NO:74;
- (j) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 65 degrees C, or 4X SSC at 42 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g); and
- (k) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 50 degrees C, or 6X SSC at 40 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g), and that has a length that is at least 25% of the length of SEQ ID NO:73.
- 83. A protein comprising an amino acid sequence selected from the group consisting of:
 - (a) the amino acid sequence of SEQ ID NO:74;
 - (b) a fragment of the amino acid sequence of SEQ ID NO:74, the fragment comprising eight contiguous amino acids of SEQ ID NO:74; and
- (c) the amino acid sequence encoded by the cDNA insert of clone vo24_1 deposited with the ATCC under accession number PTA-366; the protein being substantially free from other mammalian proteins.
- 84. An isolated polynucleotide comprising a nucleotide sequence selected from the group consisting of:

- (a) the nucleotide sequence of SEQ ID NO:75;
- (b) the nucleotide sequence of SEQ ID NO:75 from nucleotide 73 to nucleotide 798;
- (c) the nucleotide sequence of SEQ ID NO:75 from nucleotide 142 to nucleotide 798;
- (d) the nucleotide sequence of the full-length protein coding sequence of clone vo25_1 deposited with the ATCC under accession number PTA-366;
- (e) a nucleotide sequence encoding the full-length protein encoded by the cDNA insert of clone vo25_1 deposited with the ATCC under accession number PTA-366;
- (f) the nucleotide sequence of a mature protein coding sequence of clone vo25_1 deposited with the ATCC under accession number PTA-366;
- (g) a nucleotide sequence encoding a mature protein encoded by the cDNA insert of clone vo25_1 deposited with the ATCC under accession number PTA-366;
- (h) a nucleotide sequence encoding a protein comprising the amino acid sequence of SEQ ID NO:76;
- (i) a nucleotide sequence encoding a protein comprising a fragment of the amino acid sequence of SEQ ID NO:76, the fragment comprising eight contiguous amino acids of SEQ ID NO:76;
- (j) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 65 degrees C, or 4X SSC at 42 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g); and
- (k) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 50 degrees C, or 6X SSC at 40 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g), and that has a length that is at least 25% of the length of SEQ ID NO:75.
- 85. A protein comprising an amino acid sequence selected from the group consisting of:
 - (a) the amino acid sequence of SEQ ID NO:76;

(b) a fragment of the amino acid sequence of SEQ ID NO:76, the fragment comprising eight contiguous amino acids of SEQ ID NO:76; and

- (c) the amino acid sequence encoded by the cDNA insert of clone vo25_1 deposited with the ATCC under accession number PTA-366; the protein being substantially free from other mammalian proteins.
- 86. An isolated polynucleotide comprising a nucleotide sequence selected from the group consisting of:
 - (a) the nucleotide sequence of SEQ ID NO:27;
 - (b) the nucleotide sequence of SEQ ID NO:27 from nucleotide 26 to nucleotide 307;
 - (c) the nucleotide sequence of SEQ ID NO:27 from nucleotide 101 to nucleotide 307;
 - (d) the nucleotide sequence of the full-length protein coding sequence of clone vo26_1 deposited with the ATCC under accession number PTA-366;
 - (e) a nucleotide sequence encoding the full-length protein encoded by the cDNA insert of clone vo26_1 deposited with the ATCC under accession number PTA-366;
 - (f) the nucleotide sequence of a mature protein coding sequence of clone vo26_1 deposited with the ATCC under accession number PTA-366;
 - (g) a nucleotide sequence encoding a mature protein encoded by the cDNA insert of clone vo26_1 deposited with the ATCC under accession number PTA-366;
 - (h) a nucleotide sequence encoding a protein comprising the amino acid sequence of SEQ ID NO:78;
 - (i) a nucleotide sequence encoding a protein comprising a fragment of the amino acid sequence of SEQ ID NO:78, the fragment comprising eight contiguous amino acids of SEQ ID NO:78;
 - (j) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 65 degrees C, or 4X SSC at 42 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g); and

(k) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 50 degrees C, or 6X SSC at 40 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g), and that has a length that is at least 25% of the length of SEQ ID NO:27.

- 87. A protein comprising an amino acid sequence selected from the group consisting of:
 - (a) the amino acid sequence of SEQ ID NO:78;
 - (b) a fragment of the amino acid sequence of SEQ ID NO:78, the fragment comprising eight contiguous amino acids of SEQ ID NO:78; and
- (c) the amino acid sequence encoded by the cDNA insert of clone vo26_1 deposited with the ATCC under accession number PTA-366; the protein being substantially free from other mammalian proteins.
- 88. An isolated polynucleotide comprising a nucleotide sequence selected from the group consisting of:
 - (a) the nucleotide sequence of SEQ ID NO:79;
 - (b) the nucleotide sequence of SEQ ID NO:79 from nucleotide 43 to nucleotide 228;
 - (c) the nucleotide sequence of SEQ ID NO:79 from nucleotide 94 to nucleotide 228;
 - (d) the nucleotide sequence of the full-length protein coding sequence of clone vp23_1 deposited with the ATCC under accession number PTA-368;
 - (e) a nucleotide sequence encoding the full-length protein encoded by the cDNA insert of clone vp23_1 deposited with the ATCC under accession number PTA-368;
 - (f) the nucleotide sequence of a mature protein coding sequence of clone vp23_1 deposited with the ATCC under accession number PTA-368;
 - (g) a nucleotide sequence encoding a mature protein encoded by the cDNA insert of clone vp23_1 deposited with the ATCC under accession number PTA-368;

(h) a nucleotide sequence encoding a protein comprising the amino acid sequence of SEQ ID NO:80;

- (i) a nucleotide sequence encoding a protein comprising a fragment of the amino acid sequence of SEQ ID NO:80, the fragment comprising eight contiguous amino acids of SEQ ID NO:80;
- (j) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 65 degrees C, or 4X SSC at 42 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g); and
- (k) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 50 degrees C, or 6X SSC at 40 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g), and that has a length that is at least 25% of the length of SEQ ID NO:79.
- 89. A protein comprising an amino acid sequence selected from the group consisting of:
 - (a) the amino acid sequence of SEQ ID NO:80;
 - (b) a fragment of the amino acid sequence of SEQ ID NO:80, the fragment comprising eight contiguous amino acids of SEQ ID NO:80; and
- (c) the amino acid sequence encoded by the cDNA insert of clone vp23_1 deposited with the ATCC under accession number PTA-368; the protein being substantially free from other mammalian proteins.
- 90. An isolated polynucleotide comprising a nucleotide sequence selected from the group consisting of:
 - (a) the nucleotide sequence of SEQ ID NO:81;
 - (b) the nucleotide sequence of SEQ ID NO:81 from nucleotide 245 to nucleotide 427;
 - (c) the nucleotide sequence of SEQ ID NO:81 from nucleotide 308 to nucleotide 427;
 - (d) the nucleotide sequence of the full-length protein coding sequence of clone vq7_1 deposited with the ATCC under accession number PTA-368;

(e) a nucleotide sequence encoding the full-length protein encoded by the cDNA insert of clone vq7_1 deposited with the ATCC under accession number PTA-368;

- (f) the nucleotide sequence of a mature protein coding sequence of clone vq7_1 deposited with the ATCC under accession number PTA-368;
- (g) a nucleotide sequence encoding a mature protein encoded by the cDNA insert of clone vq7_1 deposited with the ATCC under accession number PTA-368;
- (h) a nucleotide sequence encoding a protein comprising the amino acid sequence of SEQ ID NO:82;
- (i) a nucleotide sequence encoding a protein comprising a fragment of the amino acid sequence of SEQ ID NO:82, the fragment comprising eight contiguous amino acids of SEQ ID NO:82;
- (j) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 65 degrees C, or 4X SSC at 42 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g); and
- (k) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 50 degrees C, or 6X SSC at 40 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g), and that has a length that is at least 25% of the length of SEQ ID NO:81.
- 91. A protein comprising an amino acid sequence selected from the group consisting of:
 - (a) the amino acid sequence of SEQ ID NO:82;
 - (b) a fragment of the amino acid sequence of SEQ ID NO:82, the fragment comprising eight contiguous amino acids of SEQ ID NO:82; and
- (c) the amino acid sequence encoded by the cDNA insert of clone vq7_1 deposited with the ATCC under accession number PTA-368; the protein being substantially free from other mammalian proteins.
- 92. An isolated polynucleotide comprising a nucleotide sequence selected from the group consisting of:

- (a) the nucleotide sequence of SEQ ID NO:83;
- (b) the nucleotide sequence of SEQ ID NO:83 from nucleotide 119 to nucleotide 475;
- (c) the nucleotide sequence of SEQ ID NO:83 from nucleotide 185 to nucleotide 475;
- (d) the nucleotide sequence of the full-length protein coding sequence of clone vq8_1 deposited with the ATCC under accession number PTA-368;
- (e) a nucleotide sequence encoding the full-length protein encoded by the cDNA insert of clone vq8_1 deposited with the ATCC under accession number PTA-368;
- (f) the nucleotide sequence of a mature protein coding sequence of clone vq8_1 deposited with the ATCC under accession number PTA-368;
- (g) a nucleotide sequence encoding a mature protein encoded by the cDNA insert of clone vq8_1 deposited with the ATCC under accession number PTA-368;
- (h) a nucleotide sequence encoding a protein comprising the amino acid sequence of SEQ ID NO:84;
- (i) a nucleotide sequence encoding a protein comprising a fragment of the amino acid sequence of SEQ ID NO:84, the fragment comprising eight contiguous amino acids of SEQ ID NO:84;
- (j) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 65 degrees C, or 4X SSC at 42 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g); and
- (k) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 50 degrees C, or 6X SSC at 40 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g), and that has a length that is at least 25% of the length of SEQ ID NO:83.
- 93. A protein comprising an amino acid sequence selected from the group consisting of:
 - (a) the amino acid sequence of SEQ ID NO:84;

(b) a fragment of the amino acid sequence of SEQ ID NO:84, the fragment comprising eight contiguous amino acids of SEQ ID NO:84; and

- (c) the amino acid sequence encoded by the cDNA insert of clone vq8_1 deposited with the ATCC under accession number PTA-368; the protein being substantially free from other mammalian proteins.
- 94. An isolated polynucleotide comprising a nucleotide sequence selected from the group consisting of:
 - (a) the nucleotide sequence of SEQ ID NO:85;
 - (b) the nucleotide sequence of SEQ ID NO:85 from nucleotide 90 to nucleotide 323;
 - (c) the nucleotide sequence of SEQ ID NO:85 from nucleotide 141 to nucleotide 323;
 - (d) the nucleotide sequence of the full-length protein coding sequence of clone vq9_1 deposited with the ATCC under accession number PTA-368;
 - (e) a nucleotide sequence encoding the full-length protein encoded by the cDNA insert of clone vq9_1 deposited with the ATCC under accession number PTA-368;
 - (f) the nucleotide sequence of a mature protein coding sequence of clone vq9_1 deposited with the ATCC under accession number PTA-368;
 - (g) a nucleotide sequence encoding a mature protein encoded by the cDNA insert of clone vq9_1 deposited with the ATCC under accession number PTA-368;
 - (h) a nucleotide sequence encoding a protein comprising the amino acid sequence of SEQ ID NO:86;
 - (i) a nucleotide sequence encoding a protein comprising a fragment of the amino acid sequence of SEQ ID NO:86, the fragment comprising eight contiguous amino acids of SEQ ID NO:86;
 - (j) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 65 degrees C, or 4X SSC at 42 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g); and

(k) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 50 degrees C, or 6X SSC at 40 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g), and that has a length that is at least 25% of the length of SEQ ID NO:85.

- 95. A protein comprising an amino acid sequence selected from the group consisting of:
 - (a) the amino acid sequence of SEQ ID NO:86;
 - (b) a fragment of the amino acid sequence of SEQ ID NO:86, the fragment comprising eight contiguous amino acids of SEQ ID NO:86; and
- (c) the amino acid sequence encoded by the cDNA insert of clone vq9_1 deposited with the ATCC under accession number PTA-368; the protein being substantially free from other mammalian proteins.
- 96. An isolated polynucleotide comprising a nucleotide sequence selected from the group consisting of:
 - (a) the nucleotide sequence of SEQ ID NO:87;
 - (b) the nucleotide sequence of SEQ ID NO:87 from nucleotide 18 to nucleotide 452;
 - (c) the nucleotide sequence of SEQ ID NO:87 from nucleotide 72 to nucleotide 452;
 - (d) the nucleotide sequence of the full-length protein coding sequence of clone vq10_1 deposited with the ATCC under accession number PTA-368;
 - (e) a nucleotide sequence encoding the full-length protein encoded by the cDNA insert of clone vq10_1 deposited with the ATCC under accession number PTA-368;
 - (f) the nucleotide sequence of a mature protein coding sequence of clone vq10_1 deposited with the ATCC under accession number PTA-368;
 - (g) a nucleotide sequence encoding a mature protein encoded by the cDNA insert of clone vq10_1 deposited with the ATCC under accession number PTA-368;

(h) a nucleotide sequence encoding a protein comprising the amino acid sequence of SEQ ID NO:88;

- (i) a nucleotide sequence encoding a protein comprising a fragment of the amino acid sequence of SEQ ID NO:88, the fragment comprising eight contiguous amino acids of SEQ ID NO:88;
- (j) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 65 degrees C, or 4X SSC at 42 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g); and
- (k) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 50 degrees C, or 6X SSC at 40 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g), and that has a length that is at least 25% of the length of SEQ ID NO:87.
- 97. A protein comprising an amino acid sequence selected from the group consisting of:
 - (a) the amino acid sequence of SEQ ID NO:88;
 - (b) a fragment of the amino acid sequence of SEQ ID NO:88, the fragment comprising eight contiguous amino acids of SEQ ID NO:88; and
- (c) the amino acid sequence encoded by the cDNA insert of clone vq10_1 deposited with the ATCC under accession number PTA-368; the protein being substantially free from other mammalian proteins.
- 98. An isolated polynucleotide comprising a nucleotide sequence selected from the group consisting of:
 - (a) the nucleotide sequence of SEQ ID NO:89;
 - (b) the nucleotide sequence of SEQ ID NO:89 from nucleotide 196 to nucleotide 378;
 - (c) the nucleotide sequence of SEQ ID NO:89 from nucleotide 262 to nucleotide 378;
 - (d) the nucleotide sequence of the full-length protein coding sequence of clone vq13_1 deposited with the ATCC under accession number PTA-368;

(e) a nucleotide sequence encoding the full-length protein encoded by the cDNA insert of clone vq13_1 deposited with the ATCC under accession number PTA-368;

- (f) the nucleotide sequence of a mature protein coding sequence of clone vq13_1 deposited with the ATCC under accession number PTA-368;
- (g) a nucleotide sequence encoding a mature protein encoded by the cDNA insert of clone vq13_1 deposited with the ATCC under accession number PTA-368;
- (h) a nucleotide sequence encoding a protein comprising the amino acid sequence of SEQ ID NO:90;
- (i) a nucleotide sequence encoding a protein comprising a fragment of the amino acid sequence of SEQ ID NO:90, the fragment comprising eight contiguous amino acids of SEQ ID NO:90;
- (j) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 65 degrees C, or 4X SSC at 42 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g); and
- (k) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 50 degrees C, or 6X SSC at 40 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g), and that has a length that is at least 25% of the length of SEQ ID NO:89.
- 99. A protein comprising an amino acid sequence selected from the group consisting of:
 - (a) the amino acid sequence of SEQ ID NO:90;
 - (b) a fragment of the amino acid sequence of SEQ ID NO:90, the fragment comprising eight contiguous amino acids of SEQ ID NO:90; and
- (c) the amino acid sequence encoded by the cDNA insert of clone vq13_1 deposited with the ATCC under accession number PTA-368; the protein being substantially free from other mammalian proteins.
- 100. An isolated polynucleotide comprising a nucleotide sequence selected from the group consisting of:

- (a) the nucleotide sequence of SEQ ID NO:91;
- (b) the nucleotide sequence of SEQ ID NO:91 from nucleotide 35 to nucleotide 718;
- (c) the nucleotide sequence of SEQ ID NO:91 from nucleotide 173 to nucleotide 718;
- (d) the nucleotide sequence of the full-length protein coding sequence of clone vq16_1 deposited with the ATCC under accession number PTA-368;
- (e) a nucleotide sequence encoding the full-length protein encoded by the cDNA insert of clone vq16_1 deposited with the ATCC under accession number PTA-368;
- (f) the nucleotide sequence of a mature protein coding sequence of clone vq16_1 deposited with the ATCC under accession number PTA-368;
- (g) a nucleotide sequence encoding a mature protein encoded by the cDNA insert of clone vq16_1 deposited with the ATCC under accession number PTA-368;
- (h) a nucleotide sequence encoding a protein comprising the amino acid sequence of SEQ ID NO:92;
- (i) a nucleotide sequence encoding a protein comprising a fragment of the amino acid sequence of SEQ ID NO:92, the fragment comprising eight contiguous amino acids of SEQ ID NO:92;
- (j) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 65 degrees C, or 4X SSC at 42 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g); and
- (k) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 50 degrees C, or 6X SSC at 40 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g), and that has a length that is at least 25% of the length of SEQ ID NO:91.
- 101. A protein comprising an amino acid sequence selected from the group consisting of:
 - (a) the amino acid sequence of SEQ ID NO:92;

(b) a fragment of the amino acid sequence of SEQ ID NO:92, the fragment comprising eight contiguous amino acids of SEQ ID NO:92; and

- (c) the amino acid sequence encoded by the cDNA insert of clone vq16_1 deposited with the ATCC under accession number PTA-368; the protein being substantially free from other mammalian proteins.
- 102. An isolated polynucleotide comprising a nucleotide sequence selected from the group consisting of:
 - (a) the nucleotide sequence of SEQ ID NO:93;
 - (b) the nucleotide sequence of SEQ ID NO:93 from nucleotide 1 to nucleotide 762;
 - (c) the nucleotide sequence of SEQ ID NO:93 from nucleotide 70 to nucleotide 762;
 - (d) the nucleotide sequence of the full-length protein coding sequence of clone vq19_1 deposited with the ATCC under accession number PTA-368;
 - (e) a nucleotide sequence encoding the full-length protein encoded by the cDNA insert of clone vq19_1 deposited with the ATCC under accession number PTA-368;
 - (f) the nucleotide sequence of a mature protein coding sequence of clone vq19_1 deposited with the ATCC under accession number PTA-368;
 - (g) a nucleotide sequence encoding a mature protein encoded by the cDNA insert of clone vq19_1 deposited with the ATCC under accession number PTA-368;
 - (h) a nucleotide sequence encoding a protein comprising the amino acid sequence of SEQ ID NO:94;
 - (i) a nucleotide sequence encoding a protein comprising a fragment of the amino acid sequence of SEQ ID NO:94, the fragment comprising eight contiguous amino acids of SEQ ID NO:94;
 - (j) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 65 degrees C, or 4X SSC at 42 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g); and

(k) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 50 degrees C, or 6X SSC at 40 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g), and that has a length that is at least 25% of the length of SEQ ID NO:93.

- 103. A protein comprising an amino acid sequence selected from the group consisting of:
 - (a) the amino acid sequence of SEQ ID NO:94;
 - (b) a fragment of the amino acid sequence of SEQ ID NO:94, the fragment comprising eight contiguous amino acids of SEQ ID NO:94; and
- (c) the amino acid sequence encoded by the cDNA insert of clone vq19_1 deposited with the ATCC under accession number PTA-368; the protein being substantially free from other mammalian proteins.
- 104. An isolated polynucleotide comprising a nucleotide sequence selected from the group consisting of:
 - (a) the nucleotide sequence of SEQ ID NO:95;
 - (b) the nucleotide sequence of SEQ ID NO:95 from nucleotide 106 to nucleotide 792;
 - (c) the nucleotide sequence of SEQ ID NO:95 from nucleotide 172 to nucleotide 792;
 - (d) the nucleotide sequence of the full-length protein coding sequence of clone vq20_1 deposited with the ATCC under accession number PTA-368;
 - (e) a nucleotide sequence encoding the full-length protein encoded by the cDNA insert of clone vq20_1 deposited with the ATCC under accession number PTA-368;
 - (f) the nucleotide sequence of a mature protein coding sequence of clone vq20_1 deposited with the ATCC under accession number PTA-368;
 - (g) a nucleotide sequence encoding a mature protein encoded by the cDNA insert of clone vq20_1 deposited with the ATCC under accession number PTA-368;

(h) a nucleotide sequence encoding a protein comprising the amino acid sequence of SEQ ID NO:96;

- (i) a nucleotide sequence encoding a protein comprising a fragment of the amino acid sequence of SEQ ID NO:96, the fragment comprising eight contiguous amino acids of SEQ ID NO:96;
- (j) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 65 degrees C, or 4X SSC at 42 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g); and
- (k) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 50 degrees C, or 6X SSC at 40 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g), and that has a length that is at least 25% of the length of SEQ ID NO:95.
- 105. A protein comprising an amino acid sequence selected from the group consisting of:
 - (a) the amino acid sequence of SEQ ID NO:96;
 - (b) a fragment of the amino acid sequence of SEQ ID NO:96, the fragment comprising eight contiguous amino acids of SEQ ID NO:96; and
- (c) the amino acid sequence encoded by the cDNA insert of clone vq20_1 deposited with the ATCC under accession number PTA-368; the protein being substantially free from other mammalian proteins.
- 106. An isolated polynucleotide comprising a nucleotide sequence selected from the group consisting of:
 - (a) the nucleotide sequence of SEQ ID NO:97;
 - (b) the nucleotide sequence of SEQ ID NO:97 from nucleotide 40 to nucleotide 315;
 - (c) the nucleotide sequence of SEQ ID NO:97 from nucleotide 124 to nucleotide 315;
 - (d) the nucleotide sequence of the full-length protein coding sequence of clone vq21_1 deposited with the ATCC under accession number PTA-368;

(e) a nucleotide sequence encoding the full-length protein encoded by the cDNA insert of clone vq21_1 deposited with the ATCC under accession number PTA-368;

- (f) the nucleotide sequence of a mature protein coding sequence of clone vq21_1 deposited with the ATCC under accession number PTA-368;
- (g) a nucleotide sequence encoding a mature protein encoded by the cDNA insert of clone vq21_1 deposited with the ATCC under accession number PTA-368;
- (h) a nucleotide sequence encoding a protein comprising the amino acid sequence of SEQ ID NO:98;
- (i) a nucleotide sequence encoding a protein comprising a fragment of the amino acid sequence of SEQ ID NO:98, the fragment comprising eight contiguous amino acids of SEQ ID NO:98;
- (j) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 65 degrees C, or 4X SSC at 42 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g); and
- (k) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 50 degrees C, or 6X SSC at 40 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g), and that has a length that is at least 25% of the length of SEQ ID NO:97.
- 107. A protein comprising an amino acid sequence selected from the group consisting of:
 - (a) the amino acid sequence of SEQ ID NO:98;
 - (b) a fragment of the amino acid sequence of SEQ ID NO:98, the fragment comprising eight contiguous amino acids of SEQ ID NO:98; and
- (c) the amino acid sequence encoded by the cDNA insert of clone vq21_1 deposited with the ATCC under accession number PTA-368; the protein being substantially free from other mammalian proteins.
- 108. An isolated polynucleotide comprising a nucleotide sequence selected from the group consisting of:

- (a) the nucleotide sequence of SEQ ID NO:99;
- (b) the nucleotide sequence of SEQ ID NO:99 from nucleotide 70 to nucleotide 699;
- (c) the nucleotide sequence of the full-length protein coding sequence of clone vr2_1 deposited with the ATCC under accession number PTA-368;
- (d) a nucleotide sequence encoding the full-length protein encoded by the cDNA insert of clone vr2_1 deposited with the ATCC under accession number PTA-368;
- (e) a nucleotide sequence encoding a protein comprising the amino acid sequence of SEQ ID NO:100;
- (f) a nucleotide sequence encoding a protein comprising a fragment of the amino acid sequence of SEQ ID NO:100, the fragment comprising eight contiguous amino acids of SEQ ID NO:100;
- (g) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 65 degrees C, or 4X SSC at 42 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(d); and
- (h) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 50 degrees C, or 6X SSC at 40 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(d), and that has a length that is at least 25% of the length of SEQ ID NO:99.
- 109. A protein comprising an amino acid sequence selected from the group consisting of:
 - (a) the amino acid sequence of SEQ ID NO:100;
 - (b) a fragment of the amino acid sequence of SEQ ID NO:100, the fragment comprising eight contiguous amino acids of SEQ ID NO:100; and
- (c) the amino acid sequence encoded by the cDNA insert of clone vr2_1 deposited with the ATCC under accession number PTA-368; the protein being substantially free from other mammalian proteins.
- 110. An isolated polynucleotide comprising a nucleotide sequence selected from the group consisting of:

- (a) the nucleotide sequence of SEQ ID NO:101;
- (b) the nucleotide sequence of SEQ ID NO:101 from nucleotide 170 to nucleotide 394;
- (c) the nucleotide sequence of SEQ ID NO:101 from nucleotide 227 to nucleotide 394;
- (d) the nucleotide sequence of the full-length protein coding sequence of clone PTA-1075 deposited with the ATCC under accession number PTA-1075;
- (e) a nucleotide sequence encoding the full-length protein encoded by the cDNA insert of clone vc69_1 deposited with the ATCC under accession number PTA-1075;
- (f) the nucleotide sequence of a mature protein coding sequence of clone vc69_1 deposited with the ATCC under accession number PTA-1075;
- (g) a nucleotide sequence encoding a mature protein encoded by the cDNA insert of clone vc69_1 deposited with the ATCC under accession number PTA-1075;
- (h) a nucleotide sequence encoding a protein comprising the amino acid sequence of SEQ ID NO:102;
- (i) a nucleotide sequence encoding a protein comprising a fragment of the amino acid sequence of SEQ ID NO:102, the fragment comprising eight contiguous amino acids of SEQ ID NO:102;
- (j) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 65 degrees C, or 4X SSC at 42 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g); and
- (k) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 50 degrees C, or 6X SSC at 40 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g), and that has a length that is at least 25% of the length of SEQ ID NO:101.
- 111. A protein comprising an amino acid sequence selected from the group consisting of:
 - (a) the amino acid sequence of SEQ ID NO:102;

(b) a fragment of the amino acid sequence of SEQ ID NO:102, the fragment comprising eight contiguous amino acids of SEQ ID NO:102; and

- (c) the amino acid sequence encoded by the cDNA insert of clone vc69_1 deposited with the ATCC under accession number PTA-1075; the protein being substantially free from other mammalian proteins.
- 112. An isolated polynucleotide comprising a nucleotide sequence selected from the group consisting of:
 - (a) the nucleotide sequence of SEQ ID NO:103:
 - (b) the nucleotide sequence of SEQ ID NO:103 from nucleotide 43 to nucleotide 198;
 - (c) the nucleotide sequence of SEQ ID NO:103 from nucleotide 85 to nucleotide 198;
 - (d) the nucleotide sequence of the full-length protein coding sequence of clone vc71_1 deposited with the ATCC under accession number PTA-1075;
 - (e) a nucleotide sequence encoding the full-length protein encoded by the cDNA insert of clone vc71_1 deposited with the ATCC under accession number PTA-1075;
 - (f) the nucleotide sequence of a mature protein coding sequence of clone vc71_1 deposited with the ATCC under accession number PTA-1075;
 - (g) a nucleotide sequence encoding a mature protein encoded by the cDNA insert of clone vc71_1 deposited with the ATCC under accession number PTA-1075;
 - (h) a nucleotide sequence encoding a protein comprising the amino acid sequence of SEQ ID NO:104;
 - (i) a nucleotide sequence encoding a protein comprising a fragment of the amino acid sequence of SEQ ID NO:104, the fragment comprising eight contiguous amino acids of SEQ ID NO:104;
 - (j) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 65 degrees C, or 4X SSC at 42 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g); and

(k) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 50 degrees C, or 6X SSC at 40 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g), and that has a length that is at least 25% of the length of SEQ ID NO:103.

- 113. A protein comprising an amino acid sequence selected from the group consisting of:
 - (a) the amino acid sequence of SEQ ID NO:104;
 - (b) a fragment of the amino acid sequence of SEQ ID NO:104, the fragment comprising eight contiguous amino acids of SEQ ID NO:104; and
- (c) the amino acid sequence encoded by the cDNA insert of clone vc71_1 deposited with the ATCC under accession number PTA-1075; the protein being substantially free from other mammalian proteins.
- 114. An isolated polynucleotide comprising a nucleotide sequence selected from the group consisting of:
 - (a) the nucleotide sequence of SEQ ID NO:105;
 - (b) the nucleotide sequence of SEQ ID NO:105 from nucleotide 260 to nucleotide 1552;
 - (c) the nucleotide sequence of SEQ ID NO:105 from nucleotide 335 to nucleotide 1552;
 - (d) the nucleotide sequence of the full-length protein coding sequence of clone vo27_1 deposited with the ATCC under accession number PTA-1075;
 - (e) a nucleotide sequence encoding the full-length protein encoded by the cDNA insert of clone vo27_1 deposited with the ATCC under accession number PTA-1075;
 - (f) the nucleotide sequence of a mature protein coding sequence of clone vo27_1 deposited with the ATCC under accession number PTA-1075;
 - (g) a nucleotide sequence encoding a mature protein encoded by the cDNA insert of clone vo27_1 deposited with the ATCC under accession number PTA-1075;

(h) a nucleotide sequence encoding a protein comprising the amino acid sequence of SEQ ID NO:106;

- (i) a nucleotide sequence encoding a protein comprising a fragment of the amino acid sequence of SEQ ID NO:106, the fragment comprising eight contiguous amino acids of SEQ ID NO:106;
- (j) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 65 degrees C, or 4X SSC at 42 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g); and
- (k) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 50 degrees C, or 6X SSC at 40 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g), and that has a length that is at least 25% of the length of SEQ ID NO:105.
- 115. A protein comprising an amino acid sequence selected from the group consisting of:
 - (a) the amino acid sequence of SEQ ID NO:106;
 - (b) a fragment of the amino acid sequence of SEQ ID NO:106, the fragment comprising eight contiguous amino acids of SEQ ID NO:106; and
- (c) the amino acid sequence encoded by the cDNA insert of clone vo27_1 deposited with the ATCC under accession number PTA-1075; the protein being substantially free from other mammalian proteins.
- 116. An isolated polynucleotide comprising a nucleotide sequence selected from the group consisting of:
 - (a) the nucleotide sequence of SEQ ID NO:107;
 - (b) the nucleotide sequence of SEQ ID NO:107 from nucleotide 15 to nucleotide 320;
 - (c) the nucleotide sequence of SEQ ID NO:107 from nucleotide 72 to nucleotide 320;
 - (d) the nucleotide sequence of the full-length protein coding sequence of clone vo31_1 deposited with the ATCC under accession number PTA-1075;

(e) a nucleotide sequence encoding the full-length protein encoded by the cDNA insert of clone vo31_1 deposited with the ATCC under accession number PTA-1075;

- (f) the nucleotide sequence of a mature protein coding sequence of clone vo31_1 deposited with the ATCC under accession number PTA-1075;
- (g) a nucleotide sequence encoding a mature protein encoded by the cDNA insert of clone vo31_1 deposited with the ATCC under accession number PTA-1075;
- (h) a nucleotide sequence encoding a protein comprising the amino acid sequence of SEQ ID NO:108;
- (i) a nucleotide sequence encoding a protein comprising a fragment of the amino acid sequence of SEQ ID NO:108, the fragment comprising eight contiguous amino acids of SEQ ID NO:108;
- (j) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 65 degrees C, or 4X SSC at 42 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g); and
- (k) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 50 degrees C, or 6X SSC at 40 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g), and that has a length that is at least 25% of the length of SEQ ID NO:107.
- 117. A protein comprising an amino acid sequence selected from the group consisting of:
 - (a) the amino acid sequence of SEQ ID NO:108;
 - (b) a fragment of the amino acid sequence of SEQ ID NO:108, the fragment comprising eight contiguous amino acids of SEQ ID NO:108; and
- (c) the amino acid sequence encoded by the cDNA insert of clone vo31_1 deposited with the ATCC under accession number PTA-1075; the protein being substantially free from other mammalian proteins.
- 118. An isolated polynucleotide comprising a nucleotide sequence selected from the group consisting of:

- (a) the nucleotide sequence of SEQ ID NO:109;
- (b) the nucleotide sequence of SEQ ID NO:109 from nucleotide 38 to nucleotide 1255;
- (c) the nucleotide sequence of SEQ ID NO:109 from nucleotide 86 to nucleotide 1255;
- (d) the nucleotide sequence of the full-length protein coding sequence of clone vo32_1 deposited with the ATCC under accession number PTA-1075;
- (e) a nucleotide sequence encoding the full-length protein encoded by the cDNA insert of clone vo32_1 deposited with the ATCC under accession number PTA-1075;
- (f) the nucleotide sequence of a mature protein coding sequence of clone vo32_1 deposited with the ATCC under accession number PTA-1075;
- (g) a nucleotide sequence encoding a mature protein encoded by the cDNA insert of clone vo32_1 deposited with the ATCC under accession number 'PTA-1075;
- (h) a nucleotide sequence encoding a protein comprising the amino acid sequence of SEQ ID NO:110;
- (i) a nucleotide sequence encoding a protein comprising a fragment of the amino acid sequence of SEQ ID NO:110, the fragment comprising eight contiguous amino acids of SEQ ID NO:110;
- (j) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 65 degrees C, or 4X SSC at 42 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g); and
- (k) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 50 degrees C, or 6X SSC at 40 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g), and that has a length that is at least 25% of the length of SEQ ID NO:109.
- 119. A protein comprising an amino acid sequence selected from the group consisting of:
 - (a) the amino acid sequence of SEQ ID NO:110;

(b) a fragment of the amino acid sequence of SEQ ID NO:110, the fragment comprising eight contiguous amino acids of SEQ ID NO:110; and

- (c) the amino acid sequence encoded by the cDNA insert of clone vo32_1 deposited with the ATCC under accession number PTA-1075; the protein being substantially free from other mammalian proteins.
- 120. An isolated polynucleotide comprising a nucleotide sequence selected from the group consisting of:
 - (a) the nucleotide sequence of SEQ ID NO:111;
 - (b) the nucleotide sequence of SEQ ID NO:111 from nucleotide 80 to nucleotide 1276;
 - (c) the nucleotide sequence of SEQ ID NO:111 from nucleotide 131 to nucleotide 1276;
 - (d) the nucleotide sequence of the full-length protein coding sequence of clone vo33_1 deposited with the ATCC under accession number PTA-1075;
 - (e) a nucleotide sequence encoding the full-length protein encoded by the cDNA insert of clone vo33_1 deposited with the ATCC under accession number PTA-1075;
 - (f) the nucleotide sequence of a mature protein coding sequence of clone vo33_1 deposited with the ATCC under accession number PTA-1075;
 - (g) a nucleotide sequence encoding a mature protein encoded by the cDNA insert of clone vo33_1 deposited with the ATCC under accession number PTA-1075;
 - (h) a nucleotide sequence encoding a protein comprising the amino acid sequence of SEQ ID NO:112;
 - (i) a nucleotide sequence encoding a protein comprising a fragment of the amino acid sequence of SEQ ID NO:112, the fragment comprising eight contiguous amino acids of SEQ ID NO:112;
 - (j) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 65 degrees C, or 4X SSC at 42 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g); and

(k) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 50 degrees C, or 6X SSC at 40 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g), and that has a length that is at least 25% of the length of SEQ ID NO:111.

- 121. A protein comprising an amino acid sequence selected from the group consisting of:
 - (a) the amino acid sequence of SEQ ID NO:112;
 - (b) a fragment of the amino acid sequence of SEQ ID NO:112, the fragment comprising eight contiguous amino acids of SEQ ID NO:112; and
- (c) the amino acid sequence encoded by the cDNA insert of clone vo33_1 deposited with the ATCC under accession number PTA-1075; the protein being substantially free from other mammalian proteins.
- 122. An isolated polynucleotide comprising a nucleotide sequence selected from the group consisting of:
 - (a) the nucleotide sequence of SEQ ID NO:113;
 - (b) the nucleotide sequence of SEQ ID NO:113 from nucleotide 202 to nucleotide 429;
 - (c) the nucleotide sequence of SEQ ID NO:113 from nucleotide 292 to nucleotide 429;
 - (d) the nucleotide sequence of the full-length protein coding sequence of clone vq23_1 deposited with the ATCC under accession number PTA-1075;
 - (e) a nucleotide sequence encoding the full-length protein encoded by the cDNA insert of clone vq23_1 deposited with the ATCC under accession number PTA-1075;
 - (f) the nucleotide sequence of a mature protein coding sequence of clone vq23_1 deposited with the ATCC under accession number PTA-1075;
 - (g) a nucleotide sequence encoding a mature protein encoded by the cDNA insert of clone vq23_1 deposited with the ATCC under accession number PTA-1075;

(h) a nucleotide sequence encoding a protein comprising the amino acid sequence of SEQ ID NO:114;

- (i) a nucleotide sequence encoding a protein comprising a fragment of the amino acid sequence of SEQ ID NO:114, the fragment comprising eight contiguous amino acids of SEQ ID NO:114;
- (j) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 65 degrees C, or 4X SSC at 42 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g); and
- (k) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 50 degrees C, or 6X SSC at 40 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g), and that has a length that is at least 25% of the length of SEQ ID NO:113.
- 123. A protein comprising an amino acid sequence selected from the group consisting of:
 - (a) the amino acid sequence of SEQ ID NO:114;
 - (b) a fragment of the amino acid sequence of SEQ ID NO:114, the fragment comprising eight contiguous amino acids of SEQ ID NO:114; and
- (c) the amino acid sequence encoded by the cDNA insert of clone vq23_1 deposited with the ATCC under accession number PTA-1075; the protein being substantially free from other mammalian proteins.
- 124. An isolated polynucleotide comprising a nucleotide sequence selected from the group consisting of:
 - (a) the nucleotide sequence of SEQ ID NO:115;
 - (b) the nucleotide sequence of SEQ ID NO:115 from nucleotide 37 to nucleotide 1113;
 - (c) the nucleotide sequence of SEQ ID NO:115 from nucleotide 88 to nucleotide 1113;
 - (d) the nucleotide sequence of the full-length protein coding sequence of clone vq24_1 deposited with the ATCC under accession number PTA-1075;

(e) a nucleotide sequence encoding the full-length protein encoded by the cDNA insert of clone vq24_1 deposited with the ATCC under accession number PTA-1075;

- (f) the nucleotide sequence of a mature protein coding sequence of clone vq24_1 deposited with the ATCC under accession number PTA-1075;
- (g) a nucleotide sequence encoding a mature protein encoded by the cDNA insert of clone vq24_1 deposited with the ATCC under accession number PTA-1075;
- (h) a nucleotide sequence encoding a protein comprising the amino acid sequence of SEQ ID NO:116;
- (i) a nucleotide sequence encoding a protein comprising a fragment of the amino acid sequence of SEQ ID NO:116, the fragment comprising eight contiguous amino acids of SEQ ID NO:116;
- (j) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 65 degrees C, or 4X SSC at 42 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g); and
- (k) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 50 degrees C, or 6X SSC at 40 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g), and that has a length that is at least 25% of the length of SEQ ID NO:115.
- 125. A protein comprising an amino acid sequence selected from the group consisting of:
 - (a) the amino acid sequence of SEQ ID NO:116;
 - (b) a fragment of the amino acid sequence of SEQ ID NO:116, the fragment comprising eight contiguous amino acids of SEQ ID NO:116; and
- (c) the amino acid sequence encoded by the cDNA insert of clone vq24_1 deposited with the ATCC under accession number PTA-1075; the protein being substantially free from other mammalian proteins.
- 126. An isolated polynucleotide comprising a nucleotide sequence selected from the group consisting of:

- (a) the nucleotide sequence of SEQ ID NO:117;
- (b) the nucleotide sequence of SEQ ID NO:117 from nucleotide 40 to nucleotide 207;
- (c) the nucleotide sequence of SEQ ID NO:117 from nucleotide 103 to nucleotide 207;
- (d) the nucleotide sequence of the full-length protein coding sequence of clone vq26_1 deposited with the ATCC under accession number PTA-1075;
- (e) a nucleotide sequence encoding the full-length protein encoded by the cDNA insert of clone vq26_1 deposited with the ATCC under accession number PTA-1075;
- (f) the nucleotide sequence of a mature protein coding sequence of clone vq26_1 deposited with the ATCC under accession number PTA-1075;
- (g) a nucleotide sequence encoding a mature protein encoded by the cDNA insert of clone vq26_1 deposited with the ATCC under accession number PTA-1075;
- (h) a nucleotide sequence encoding a protein comprising the amino acid sequence of SEQ ID NO:118;
- (i) a nucleotide sequence encoding a protein comprising a fragment of the amino acid sequence of SEQ ID NO:118, the fragment comprising eight contiguous amino acids of SEQ ID NO:118;
- (j) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 65 degrees C, or 4X SSC at 42 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g); and
- (k) the nucleotide sequence of a polynucleotide that hybridizes under conditions at least as stringent as 4X SSC at 50 degrees C, or 6X SSC at 40 degrees C with 50% formamide, to any one of the polynucleotides specified by (a)-(g), and that has a length that is at least 25% of the length of SEQ ID NO:117.
- 127. A protein comprising an amino acid sequence selected from the group consisting of:
 - (a) the amino acid sequence of SEQ ID NO:118;

there will also be a server of the confidence of

(b) a fragment of the amino acid sequence of SEQ ID NO:118, the fragment comprising eight contiguous amino acids of SEQ ID NO:118; and

(c) the amino acid sequence encoded by the cDNA insert of clone vq26_1 deposited with the ATCC under accession number PTA-1075; the protein being substantially free from other mammalian proteins.